AA505075-derived oligonucleotide SEQ ID 4947.

```
Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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31-OCT-2002

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Ś Li Y, Sa , Tang L, Sandrasagra A, K L, Shahabuddin S; Katz E, Pabalan J, Aguilar Ö

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent. ខ

SEQ ID NO 4947; 763pp; English.

CC surfactant depletion or hyposecretion, when administered to a mammal. The cappression of a target polypeptide associated with lung airway or lung construction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition confidence of the invention has antiallergic, antiinflammatory, antiasthmatic, and the invention has antiallergic, antiinflammatory, antiasthmatic, and the invention has antiallergic, antiinflammatory, antiasthmatic, confidence of the invention of the composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction and/or lung confilammation, allergies, asthma, impeded respiration, respiratory confilammation, allergies, asthma, impeded respiration, respiratory confilammation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the middless present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system content of the arget RNA serves to prevent the breakdown of the product of free oligonucleotides into products that free adenosine into the system content of the serves to prevent the breakdown of the product of the content of the system content of the serves to prevent the breakdown of the product of the content of the serves to prevent the breakdown of the product of the content of the serves to prevent the breakdown of the product of the content of the serves to prevent the breakdown of the content of the serves to prevent the breakdown of the conten This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors. unwanted effects due to it (A) receptors,

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Query Match Best Local ! Matches Local 16; Similarity Conservative 1.1%; 0; Mismatches Score 14.8; Pred. No. 4 4.4e+02 Length Indels 20 <u>.</u> Gaps

0

18

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ABD25936 standard; DNA;

ABD25936;

(first entry)

AA505075-derived oligonucleotide SEQ ID 4948.

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-drenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory discress syndrome; allergic rhinitis; pulmonary hypertension; respiratory discress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; pulmonary transplantation rejection; ss; primer. cancer; bronchitis;

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Nyce JW, Miller S, Ś Li Y, Sandrasagra A, L, Shahabuddin S Katz ū Pabalan Aguilar

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WPI; 2003-093058/08

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, target nucleic acids associated with lung airway or lung dysfunction, bronchodilating agent. targeted ទ

Claim 15; SEQ ID NO 4948; 763pp; English.

RESULT 752
ABD25936
ID ABD259
XX ABD259
XX ABD259
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XX ABD259
XX ABD259
XX AA5050
XX Human;
XX Feepir
XX Emphron
XX Human;
XX Feepir
XX Fe CC comprising oligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and composition gadenosine sensitivity, levels of adenosine (A) or (A) receptors, compression of a target polyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating compression of a target polyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating compression of a target polypositide associated with lung airway or lung composition or cancer and can be anti-sense to the corresponding mRNA. The composition is experted to reduce the invention also describes a kit, that comprises: (a) a delivery composition of the invention has antiallergic, antiinflammatory, antiasthmatic, compression, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypoptide present in the lungs. The composition such as pulmonary vasoconstriction, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, comprises of pain, cystic fibrosis, allergic rininitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary typertension, emphysema, chronic obstructive pulmonary disease, pulmonary composition for the contract of the articles of context. transplantation rejection, pulmonary infections, bronce The reduced adenosine content of the anti-sense oligos This invention describes a novel composition (a) a first active agent,

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Best Local S
Matches 16
                         comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
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Miller S,
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the oligonucleotides into pro
e.g., lung, brain, heart, kid
prevent any unwanted effects
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                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel composition (a) a first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 15;
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L, Shahabuddin
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                                                                                                                                                        Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targe nucleic acids associated with lung airway or lung dysfunction, bronchodilating agent.
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Tang L,
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                                                                                                                                                                                                                                                                                                                                                                                              Sandrasagra A,
L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                              Katz E,
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Pabalan
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                                                                                                                                                                                      nosine, targeted to dysfunction, and
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Claim

15;

SEQ ID NO

553;

763pp; English.

comprising This invention

oligonucleotides,

describes

a novel composition (a) a fir des, effective for alleviating

first

active agent,

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cc bronchoconstriction, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, consumers of the ministered to a mammal. The colligonucleotides are derived from a gene encoding or regulating content of a target polypeptide associated with lung airway or lung dysfunction of a target polypeptide associated with lung airway or lung content on also describes a kit, that comprises: (a) a delivery construction also describes a kit, that comprises: (b) the oligonucleotides, (c) device, in separate containers, (b) the oligonucleotides, (c) consists of the invention has antiallergic, antiinflammatory, antiashmatic, analysesic, hypotensive, immunosuppressive and cytostatic activity, is a composition comprises oligo and is administered to reduce the production composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to revulve the amount of target polypeptide present in the lungs. The composition, allergies and/or burdaction and/or lung conflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, respiratory conflammation, allergies, asthma, impeded respirator, respiratory conflammation, allergies, chronic obstructive pulmonary disease, into norduct anti-sense oligos corresponding to the oligonucleotion in the target RNA serves to prevent the breakdown of the surfacent in the target RNA serves to prevent the breakdown of the ministers into the target RNA serves to prevent the breakdown of the ministers into the serves to prevent the breakdown of the ministers and notice in the target RNA serves to prevent the breakdown of the ministers and notice and adenosine into the analyse and notice and notice and and notice in the breakdown of the ministers.
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  Nyce JW,
Miller S,
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                                                                                                                                24-APR-2001; 2001US-0286036P
                                                                                                                                                                                  23-APR-2002; 2002WO-US013143.
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16; Conserv
                                                                             EPIGENESIS
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Tang
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ilarity 88.9%;
Conservative
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  Sandrasagra A,
L, Shahabuddin
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                                                                                PHARM INC.
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Pred. No. 4.4e
O; Mismatches
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                             Pabalan
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cc comprising oligonucleotides, effective for alleviating componentation, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, csurfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating cypression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung cypression of a target polypeptide associated with lung airway or lung expression of a target polypeptide associated with lung airway or lung cypression of the invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition of the invention as antiallergic, antiantiallergic, antiatively, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The cypressise and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies, asthma, impeded respiration, respiratory changes reduced adenosine content of the anti-sense oligos corresponding to the millione present in the target RNA serves to prevent the breakdown of the manual products that free adenosine into the system content of the anti-sense oligos corresponding to the oligonacional and thereby, to prevent any unwanted effects due to it
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Best Local 9
                                                    Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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                                                                                                                                                                                                                                                   Human stanniocalcin-derived oligo SEQ
                                                                                                                                                                                                                                                                                                                                    ABD21765;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 15; SEQ ID NO 4683; 763pp; English.
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                                                                                                                                                                                                                                                                                                                                                       CC surfactant depletion or hyposecretion, when administered to a mammal. The CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.

CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or composition comprises oligo and is administered to reduce the production composition, allergies and/or bronchoconstriction and/or lung creating a respiratory, lung or malignant disease. The administered composition is amount of target polypeptide present in the lungs. The cc pulmonary obstruction, and/or bronchoconstriction and/or lung cinflammation, allergies and/or bronchoconstriction and/or lung cinflammation, allergies, asthma, impeded respiration, respiratory composition such as pulmonary vasoconstriction, pulmonary changes and condition such as pulmonary vasoconstriction are associated conflammation, emphysema, chronic obstructive pulmonary disease, pulmonary cranges and conditions, pain, cystic fibrosis, allergic rhinitis, pulmonary cranges and conditions, pronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonary brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                RESULT 757
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Best Local Similarity
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 AA281534-derived oligonucleotide SEQ ID 3687.
                                      29-JUL-2004
                                                                          ABD24675;
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                                                                                                                                                                                                                                                                                                                                                       G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                 Score 14.8;
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716 TTCTCTGTTTTGCTGTTG

Query Match Best Local : Matches 1:

Similarity

1.1%;

Score 14.8; Pred. No. 4

3; DB 1; 4.4e+02;

Length 20; Indels

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Conservative

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CC omprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating contracts and can be anti-sense to the corresponding mRNA. The invention or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery construction or cancer and can be anti-sense to the corresponding mRNA. The instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, and persent to instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, and persent to it and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or creating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascoonstriction.

CC inflammation, allergies, asthma, impeded respiration, respiratory conditions under the surfactant hypoproduction are associated with a cisease or condition such as pulmonary vascoonstriction and/or lung inflammation, allergies, asthma, impeded respiration, respiratory conditions on the surfactant hypoproduction are associated transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the minimum the b
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                                                the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.
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TTCTCTGTTCTGGTGTTG 19

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CC comprising oligonucleotides, effective for alleviating controlled adenosine sensitivity, levels of adenosine (A) receptors, cc surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating content of a target polyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating content or cancer and can be anti-sense to the corresponding mRNA. CC instruction also describes a kit, that comprises: (a) a delivery construction for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a cc treating a respiratory, lung or malignant disease. The administered composition composition composition composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypoptide present in the lungs. The composition and adsease or condition such as pulmonary vasoconstriction are associated with a disease or condition such as pulmonary vasoconstriction, respiratory constructive pulmonary disease, pulmonary disease, pulmonary disease, pulmonary disease, pulmonary cransplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of composition of the target not the breakdown of compositions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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L, Shahabuddin
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This invention describes a novel composition (a) a first active agent, CC comprising oligonucleotides, effective for alleviating CC comprising oligonucleotides, effective for alleviating CC bronchoconstriction, respiratory tract inflammation, allergies and CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The CC oligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung alivery or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) constructions for adding a carrier and for use of the kit. The composition CC of the invention has antiallergic, antinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered
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Matches 16
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Pred. No. 4
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RESULT 76
ABD22850
ID ABD22850
AC ABD2
XX ABD2
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KW respl
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergio; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                    Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                               Claim 15; SEQ
                                                                                                                      bronchodilating agent
                                                                                                                                                                                                                                                              Miller
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                                                                                                                                                                                                                                                          Li Y, Sa
Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAAAAAAAAAAAAAAA 18
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                                                                               ID NO 1862; 763pp; English
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L, Shahabuddin
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Pred. No. 4.4e+02
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This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and

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Best Local S
Matches 16
                  Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                              Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic,
                                                                                                                    24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                  31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AI652901-derived oligonucleotide SEQ ID 3508.
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                                      Sandrasagra A,
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                    Shahabuddin
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Pred. No. 4.4e
0; Mismatches
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                                        Katz E,
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RESULT 762
ABD25532
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Best Local :
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                                                                                                                                 Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AI125651-derived oligonucleotide SEQ ID 4544.
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   WO200285309-A2
                                                                      Homo sapiens.
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                             CC comprising oligonucleotides, effective for alleviating
CC producing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC oligonucleotides are derived from a gene encoding or regulating
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (A) a delivery
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
CC treating a respiratory, lung or malignant disease. The administered
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC reduce the amount of target polypeptide present in the lungs. The
CC pulmonary obstruction, and/or bronchoconstriction and/or lung
CC with a disease or condition such as pulmonary vascoonstriction
CC inflammation, allergies, asthma, impeded respiration, respiratory
CC the reduced adenosine content of the anti-sense oligos corresponding to
CC the oligonucleotides into products RNA serves to prevent the breakdown of
CC the oligonucleotides into products that free adenosine into the system
CC the oligonucleotides into products that free adenosine into the system
CC prevent any unwanted effects due to it
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                                            29-JUL-2004
                                                                             ABD25046
                                                                                                             ABD25046 standard; DNA; 20
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                                                                                                                                                                                                                                                            1 Similarity
16; Conserv
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                                                                                                                                                                                                                                                                                                                             BP;
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AI128305-derived oligonucleotide

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cc surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating compression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery comparison for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, comparises; (b) the oligonucleotides, (c) the invention has antiallergic, antiinflammatory, antiasthmatic, is a composition compressed and cytostatic activity, is a composition composition composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition composition composition is useful for preventing or reduce the amount of target polypeptide present in the production composition, allergies and/or surfactant hypoproduction are associated with a disease or condition such as gulmonary vasconstriction, inflammation, allergies, asthma, impeded respiration, respiratory transplantation rejection, pulmonary infections, bronchitis or cancer. CC inflammation, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. CC thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system ce.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                               Query Match
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CC composition of ligonucleotides, effective for alleviating composition of ligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, composition or hyposecretion, when administered to a mammal. The CC surfactant depletion or hyposecretion, when administered to a mammal. The CC oligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a cc treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The CC unifiammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary version and associated conflammation, allergies, asthma, impeded respiration, respiratory conditions polyperation, pulmonary disease, pulmonary disease, pulmonary conditions present in the target RNA serves to prevent the breakdown of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the anti-sense oligos corresponding to the charget mind the target RNA serves to prevent the breakdown of the sentement of the anti-sense oligos corresponding to the sentement in the target RNA serves to prevent the breakdown of the sentement of the anti-sense oligos corresponding to the sentement.
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RESULT 766
ABD23911/c
ID ABD239
XX ABD239
AC ABD233
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XX Human
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Best I
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Miller S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hyportension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
  comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies reducing adenosine sensitivity, levels of adenosine (A) or (A)
                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABD23911;
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                                                                                   This invention describes a novel composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-APR-2002; 2002WO-US013143.
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                                                                                                                                                                                                                                                                                                                                                                                             Sandrasagra A,
L, Shahabuddin
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Pred. No. 4
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S;
                                                                                                                                          English.
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                                                                                   (a) a first active agent,
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receptors,

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RESULT 767
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Best Local (
                                              Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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    WPI; 2003-093058/08
                                                                                                                                                                   24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                               23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                               31-OCT-2002.
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les 16; Conserv
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                                                                                                                       EPIGENESIS PHARM INC.
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C, Shahabuddin
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Pred. No. 4.4e+02;
0; Mismatches 2
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Comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising adenosine sensitivity, levels of adenosine (A) or (A) receptors, controlled the sensitivity, levels of adenosine (A) or (A) receptors, controlled the sensitivity, levels of adenosine (A) or (A) receptors, controlled the sensitivity, levels of adenosine (A) or (A) receptors, controlled the sensitivity, when administered to a mammal. The colligonucleotides are derived from a gene encoding or regulating controlled the corresponding mRNA.

Controlled the sensitivity or lung allevies (A) a delivery controlled the controlled the corresponding mRNA.

Controlled the invention also describes a kit, that comprises (A) a delivery controlled the invention and sentiallergic, antiinflammatory, antiasthmatic, confidence invention has antiallergic, antiinflammatory, antiasthmatic, controlled the invention has antiallergic, antiinflammatory, antiasthmatic, controlled the invention of the composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung ulmonary obstruction, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasconstriction, respiratory hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary continued adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content and effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targe nucleic acids associated with lung airway or lung dysfunction,
  Sequence 20
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                                                  lung, brain,
nt any unwant
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  BP;
                                                  brain, heart, kid
20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                  due to
                                                  etc, tissue environment and to it
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                                 Matches
                                         Query Match
Best Local (
                1520 AAAAAAAAAAAAGTAAAA 1537
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1 AAAAAAAAAAAAAAAAAA
                                          Similarity
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                                         1.1%;
 18
                                 <u>,</u>
                                         Score 14.8;
Pred. No. 4
                                  Mismatches
                                          4.4e+02
                                                   DB 1;
                                 2
                                                  Length 20;
                                 0
                                 Gaps
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Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                              AI125228-derived
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Homo sapiens.

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RESULT 769
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ID ADH086
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AC ADH086
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DT 11-MAR
XX
DE Nanote
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CHINCIP
                                                                                                                                                                                                                                                                                                                                                                                                                                            cc reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, cc oligonucleotides are derived from a gene encoding or regulating cc expression of a target polypeptide associated with lung airway or lung cystuction or cancer and can be anti-sense to the corresponding mRNA. Cc The invention also describes a kit, that comprises: (a) a delivery cd device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, cc instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, cc instruction for agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The cc pulmonary obstruction, and/or bronchoconstriction of the target mRNA or to inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascoonstriction are associated with a disease or condition such as pulmonary vascoonstriction are associated with a disease or condition such as pulmonary vascoonstriction are associated with a disease or condition such as pulmonary vascoonstriction are associated with a disease or condition such as pulmonary vascoonstriction are associated with a disease, pain, cystic fibrosis, allergic rhinitis, pulmonary theorem of the reduced adenosine content of the anti-sense oligos corresponding to the middless present in the target RNA serves to prevent the breakdown of the content of the anti-sense oligos corresponding to the content of the system of the content of the system of the system of the content of the system of the content of the system of the content of the system of the co
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allerg reducing adenosine sensitivity, levels of adenosine (A) or
  Linking oligonucleotide; 88; nucleic acid detection;
                                         Nanotechnology nucleic acid detection method associated
                                                                                      11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 15; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic solds associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nyce JW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-OCT-2002
                                                                                                                                                                    ADH08684 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes a novel composition (a) a first active agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                                                                   1520 AAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                            16;
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, Tang L,
                                                                                                                                                                                                                                                                          AAAAAAAAAAAAAAAA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 20
                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ID NO 4123; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                            A; 0 C; 0 G; 0 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                           1.1%;
                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                              Score 14.8;
Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                            0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pabalan J,
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د
                                                                                                                                                                                                                                                                                                                                                                                                 Length 20;
                                                                                                                                                                                                                                                                                                                                                            Indels
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                                             #54.
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                                                                                                                                                                                                                                                                                                                                                            0
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RESULT 770
ADH08814
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                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method of detecting a nucleic acid with at least two portions by providing a type of nanoparticle-oligonucleotide conjugate, contacting the nucleic acid and nanoparticles to allow hybridisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by hybridisation. The oligonucleotides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. This sequence represents a linking oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
26-APR-2000;
26-JUN-2000;
                                                                                                  Linking oligonucleotide; 88; nucleic acid detection; nanoparticle-oligonucleotide conjugate.
                                                                                                                                          Nanotechnology nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting nucleic acid used for e.g. diagnosis of diseases, fo DNA sequencing, comprises observing detectable change caused by hybridization of nucleic acid with substrate or particle bound
12-OCT-2001; 2001US-00976617.
                           26-SEP-2002.
                                                    US2002137072-A1
                                                                             Synthetic
                                                                                                                                                                      11-MAR-2004
                                                                                                                                                                                                ADH08814;
                                                                                                                                                                                                                       ADH08814 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 18; SEQ ID NO 55; 130pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-059018/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-OCT-2001; 2001US-00973638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2002137070-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nanoparticle-oligonucleotide conjugate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tacon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NANO-)
                                                                                                                                                                                                                                                                                                                1520 AAAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                            BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Letsinger RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-00344667
2000US-0200161P
2000US-00603830
                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96US-0031809P
97WO-US012783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-00240755
                                                                                                                                                                                                                                                                                                                                                      1.1%;
                                                                                                                                           acid detection method associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mucic RC,
                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                  Score 14.8; DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                      Pred. No.
                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                      4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Elghanian R;
                                                                                                                                            #54.
                                                                                                                                                                                                                                                                                                                                           0
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RESULT 771
ADH0874
ID ADH087
XX ADH087
XX ADH087
XX I1-MAR
XX Linkin
KW Linkin
KW Inanopa
XX Synthe
XX US2002
XX US2002
PD 26-SEE
XX 29-JUI
PR 29-JUI
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Best Local S
Matches 16
                                                                     29-JUL-1996;
21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
26-APR-2000;
26-JUN-2000;
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21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
26-APR-2000;
26-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of detecting a nucleic acid with at least two portions by providing a type of nanoparticle-oligonucleotide conjugate, contacting the nucleic acid and nanoparticles to allow hybridisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by hybridisation. The oligonucleotides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. This sequence represents a linking oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                         10-OCT-2001; 2001US-00974007.
                                                                                                                                                                                                                                                                                                                                                                                                  26-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2002137071-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Linking oligonucleotide; ss; nanoparticle-oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nanotechnology nucleic acid detection method associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADH08749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH08749 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oligonucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-059020/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mirkin
       (NANO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NANO-) NANOSPHERE INC.
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16; Conserv
          NANOSPHERE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ААААААААААСТААА 1537
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; 96US-0031809P.
; 97WO-US012783.
; 99US-00240755.
; 99US-00344667.
; 2000US-0200161P.
; 2000US-00603830.
                                                                     96US-0031809P.
97WO-US012783.
99US-00240755.
99US-00344667.
2000US-0200161P.
2000US-00603830.
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llarity 88.9%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid detection; conjugate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8;
Pred. No. 4.
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Mismatches
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U;
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4.4e+02;
2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Elghanian
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RESULT 772
ADH65941/c
ID ADH659
XX ADH659
XX ADH659
XX ADH659
XX Human
XX Antise
KW Inflam
KW Cardic
KW Phosph
XX Homo g
XX HOMO g
XX HOMO g
YX Claim
XX Claim
XX Claim
XX Claim
XX Claim
XX Claim
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a method of detecting a nucleic acid with at least two portions by providing a type of nanoparticle-oligonucleotide conjugate, contacting the nucleic acid and nanoparticles to allow hybridisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by hybridisation. The oligonucleotides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. This sequence represents a linking oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                       antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting nucleic acid used for e.g. diagnosis of diseases, forensics and DNA sequencing, comprises observing detectable change caused by hybridization of nucleic acid with substrate or particle bound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0
                                                                                                                                                       Crosby SD,
                                                                                                                                                                                                                  20-MAY-2002; 2002US-0381857P.
                                                                                                                                                                                                                                                 20-MAY-2003; 2003WO-US016084.
                                                                                                                                                                                                                                                                               04-DEC-2003.
                                                                                                                                                                                                                                                                                                            WO2003099215-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH65941;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADH65941 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-059019/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Taton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mirkin CA,
                                                                                                                                                                                      (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                          2004-035034/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  glucocorticoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     щ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAAAAAAAAAAAAAA 18
                                                                                                                                                       Nalseth AE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Letsinger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptor-specific antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                55; 130pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mucic RC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.4e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Elghanian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, ob

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Cushing's

Claim

4; SEQ

IJ

985pp; English. hyperlipidemia

disorder, NO 2775;

comprises

an

antisense oligonucleotides

that

are

targeted

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RESULT 773
ADH67409/c
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Best Local S
 Matches
                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothicate backbone.
                                                                                                                            The invention comprises an antisense oligonucleotides that are targeted to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obseity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets
                                                                                                                                                                                                                                                                                             New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia, Cushing's syndrome; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                                                Sequence
                                                                                               the human glucocorticoid contains 2'-methoxyethyl
                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 4243; 985pp; English.
                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-035034/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-MAY-2002; 2002US-0381857P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAY-2003; 2003WO-US016084.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003099215-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense oligonucleotide; glucocorticoid receptor; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human glucocorticoid receptor-specific antisense oligonucleotide #4243.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADH67409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                 (PHAA ) PHARMACIA CORP.
Local Similarity
nes 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1522 AAAAAAAAAAAGTAAAAGG 1539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16;
                                                                                                                                                                                                                                                                                                                                                                                                  SD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAAGAAAAAAATAAAAGG 2
                                                                ₿P;
                                                                                                                                                                                                                                                                                                                                                                                                  Nalseth AE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 C; 1 G; 15 T; 0 U; 0 Other;
                                                                1 C; 1 G; 17 T; 0 U; 0 Other;
               88.9%;
                              1.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                               receptor gene. NOTE: The present seque (2'-MOE) wings and a phosphorothicate
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
             Score 14.8; DB 1; Length 20; Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1; Length Pred. No. 4.4e+02;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                               The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20;
<u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human;
                                                                                                 backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
Gaps
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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RESULT 775 ADI47212

ADI47212 standard; DNA;

20 BP

HXXXX

ADI47212;

22-APR-2004

(first entry)

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RESULT 774
ADI34492
ID ADI344
XX ADI344
XX Nucleo
XX Nuclei
XX Nuclei
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX HOZO03
XX II-DEC
XX ADI344
XX Synthe
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                                                                                                                                                                                                                                                                                                                                                                 The invention relates to amplifying for RNA in a sample comprises comparises completed involves synthesizing single-stranded cDNA by concluding the method involves synthesizing single-stranded cDNA by converting the sample RNA with reverse transcriptase and an coligonucleotide primer that primes synthesis in a direction toward 5' end coligonucleotide primer that primes synthesis in a direction toward 5' end coligonucleotide primer that primes synthesis in a direction toward 5' end coligonucleotide from the transcription sample; and conform a transcription sample containing a cDNA template; and conform a transcription template into RNA using an RNA polymerase. The method is useful for improving RNA polymerase based RNA transcription conform a polynucleotide template. The method inhibits the undesired non-conform a polynucleotide template. The method inhibits the undesired non-conformate derived production of RNA in the transcription reaction. The present sequence represents an oligonucleotide used to exemplify RNA collection in the presence of single- and double-stranded
                                                                                                                                                                  Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amplifying for RNA in a sample, useful for improving RNA polymerase based RNA transcription from a polynucleotide template, comprises eliminating single-stranded oligonucleotide from the transcription sample.
                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-MAY-2002; 2002US-0384454P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-MAY-2003; 2003WO-US017103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-DEC-2003
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                                                                                                                                                                                                                                                                                                                                               oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; SEQ ID NO 11; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kamme FC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003102243-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleotide sequence of a dA20 oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADI34492 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (JANC ) JANSSEN PHARM NV
                                                                    1520 AAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1521 АЛАЛАЛАЛАЛАСТАЛАД 1538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
                                                                                                                                         16;
1 AAAAAAAAAAAAAAAAAA 18
                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zhu JY;
                                                                                                                                                                                                                                                                           B₽;
                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplification; RNA transcription; RNA polymerase; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                           20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                  1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ₽₽
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                                                                                                                                     Score 14.8; DB 1
Pred. No. 4.4e+02
0; Mismatches
                                                                                                                                                                                                             DB 1;
                                                                                                                                         <u>.</u>
                                                                                                                                         Gaps
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RESULT 776
ADJ51142/c
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel method involving forming a laminar flow, by passing into a micro channel, a solution containing the specimen molecules, and a solution containing probe molecules capable of forming a complex with the specimen molecules. The dispersion of the formed complex is selectively promoted, based on their affinity, and the degree of dispersion of the complex formed between the specimen molecules and the probe molecules is detected and analysed. The probe molecules are capable of producing fluorescence. This polynucleotide sequence represents an oligo used in the exemplification of the invention.
                                                                              modified_base
                                                                                                                       Synthetic
                                                                                                                                                                                        Polyalkyleneamine-conjugated oligonucleotide #1.
                                                                                                                                                                                                                      06-MAY-2004
                                                                                                                                                                                                                                                ADJ51142;
                                                                                                                                                                                                                                                                         ADJ51142
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 9; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 forming molecule containing solution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Analysis of sample molecules such as DNA fragment, by using micro channel to form laminar flow of specimen molecule-containing solution and complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-180318/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yamashita K,
Yamaguchi Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-JUL-2002; 2002JP-00211462.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-JUL-2003; 2003WO-JP009142.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       laminar flow; micro channel; complex; selectively promoted; fluorescence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecule analysing
                                                                                                                                                  inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NAAD-)
                                                                                                                                                                Antimicrobial; Antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                        1520
                                                                                                                                                                                                                                                                                                                                               H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAT
                                                                                                                                                                                                                                                                            standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                               AAAAAAAAAAAAAAAA 18
                                                                                                                                                                                                                                                                                                                                                                       АААААААААААСТАААА 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INST
                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maeda H,
                                                                                                                                                   tumour.
           //mod_base= OTHER
//mod= "Optionally conjugated with spermine,
/note= "Optionally conjugated with spermine,
polyethylenimine (PEI) 600 or PEI 1200,
tetraethylenpentamine. Also optionally 5'-pr
                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADVANCED IND SCI & TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                               1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   microchannel method related probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shimizu H,
                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 1;
Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Miyazaki M,
                                                                                                                                                                Cytostatic; prodrug; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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            5'-protected
                                                                                                                                                                                                                                                                                                                                                                                                                              20
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RESULT 777
ADX98410/c
ID ADX994
AC ADX994
AC ADX994
XX O6-MAY
XX DE Primer
XX human;
XX human;
XX human;
XX Synthe
XX JP2003
XX
                                                                                                                                                                                                                                                                                                                                                       S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              fusogenic part is a lipophilic polyamine, polyethylenimine, polyalylamine, fusogenic peptide, oligomeric imidazole, histidine, pyridine, hydroxylamine, substituted hydroxylamine, hydroxylamine, pyridine, pyridine, hydrazine, thiourea or imine. The targeting part is a ligand that binds to a cellular reporter, where the targeting part is transferrin, folate, epidermal growth factor, nerve growth factor, insulin, alpha-fetoprotein, galactose, galactosamine, lactose, mannose, a polyclonal antibody, monoclonal antibody, vitamin Bl2, ibuprofen, cholesterol, low-density lipoprotein, peptide comprising an arginine-collectine-aspartic acid sequence. The oligomeric part is an oligonucleotide, and oligonucleotide analogue, a peptide nucleic acid or a peptide nucleic acid analogue. OC is useful as a prodrug, useful in captide mucleic acid analogue, a peptide nucleic acid or a peptide nucleic acid analogue, of inserving or delaying infection, inflammation or tumour formation in organisms. The present sequence represents an oligonucleotide used in the
                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      part, a fusogenic part, and a targeting part; and enhancing the cellular uptake of OC, by conjugating OC to a fusogenic part. In (C), the fusogenic part is covalently linked to the oligomeric part. The targeting part is covalently linked to the oligomeric or fusogenic part, where the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel polyalkyleneamine-containing oligomeric compound useful for preventing or delaying infection, inflammation or tumor formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a polyalkyleneamine-containing oligomeric compound (OC). Also described is a compound (C) comprising an oligomeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mancharan M, Guzaev AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JUL-2002; 2002US-00199585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2004019000-A1
                    JP2003259875-A.
                                                   Synthetic
                                                                                      human; single nucleotide polymorphism;
                                                                                                                           Primer of the
                                                                                                                                                             06-MAY-2004
                                                                                                                                                                                                ADK98410;
                                                                                                                                                                                                                                   ADK98410 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3; Page 22; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MANO/) MANOHARAN M. (GUZA/) GUZAEV A P. (MAIE/) MAIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUL-2002; 2002US-00199585
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                                                                                                                                                                                                                                                                                                                        20 AAAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the invention.
                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                             (first
                                                                                                                           invention #4130.
                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                88.9%;
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                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. 4.4e+02
                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                          SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                          88;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20
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16-SEP-2003

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ARESULT 778
ADJ60935/c
ID ADJ609
XX ADJ609
XX O6-MAY
XX O1igon
XX Interl
XW interl
XW cystic
XX Homo s
XX 
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Best Local Similarity
Matches 16; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polynorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polynorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome;
                                                                             Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCRI, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                 WPI; 2004-203534/19.
                                                                                                                                                                                                                                                                     Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                 29-JUL-2002; 2002US-0399076P
                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-JUL-2003; 2003WO-US023509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004011613-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide associated to PDE4C #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleotide polymorphism in human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-093977/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                                  (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-MAR-2002; 2002JP-00064373
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 899
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 7439; 2627pp; Japanese.
                                                                                                                                                                                                                                                                Tang L,
lin S, Lu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTCACCTCTGAGGCCTCT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                Sandrasagra
H, Cong H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lung
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8;
Pred. No. 4
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                                                                                                                                                                                                                                                                                           Aguilar D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.4e+02;
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                                                                                                                                                                                                                                                                                              Miller
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                                                                                                                              e.g.,
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SEQ

ID NO 1791; 85pp; English

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RESULT 779
AD132920
ID AD1329
XX AD1329
XX AD1329
XX AD1329
XX Oligo
XX Innop
XX Inno
XX Innop
XX Inno
XX Inn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                              28-MAR-2000;
26-APR-2000;
26-JUN-2000;
26-JUN-2000;
11-AUG-2000;
                                                                                                                                                     09-APR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
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                                                                                                                                                                                                                                                                               11-DEC-
12-JAN-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nanoparticle; cell line auth
   (PARK/)
(TATO/)
(MIRK/)
                                                                                                                                                                                                                                               28-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                        08-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                     08-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      obstruction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3-JAN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1520 АЛАЛАЛАЛАЛАГТАЛА 1537
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                                                                                                                                                                                                                                                                        -2000;
-2001;
) PARK S.
) TATON T A.
) MIRKIN C A.
                                                                                                                                                                                                                                                                                                                                        -2000;
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authentication; gene therapy; ss; gold colloid conjugate
                                                                                                                      2000US-0176409P

2000US-012699P

2000US-0200161P

2000US-02001390

2000US-0224631P

2000US-0254392P

2000US-025431SP

2000US-0255235P

2000US-0255236P

2000US-0255236P

2001US-00820279

2001US-0282640P

2001US-0282640P

2001US-0282640P

2001US-03278640P

2001US-03278640P

2001US-03278640P

2001US-03278640P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002US-00266983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-00240755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97WO-US012783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-0031809P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present sequence represents an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1;
Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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RESULT 780
AD132905
ID 329
XX AD1329
XX AD1329
XX O6-MAY
XX O7
DE Synthe
XX nanopa
KW cell l
KW probe.
XX O5-NOV
XX O6-NOV
XX O8-OC1
XX 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel method for detecting a nucleic acid having at least two portions comprising contacting the nucleic acid with at least two types of nanoparticles, such as gold, having attached oligonucleotides and observing a detectable change brought about by hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid. The method of the invention may be useful for detecting a nucleic acid, preferably viral RNA or DNA, bacterial DNA, a gene associated with a disease, a fungal DNA, synthetic DNA or RNA, cor structurally modified natural or synthetic DNA or RNA or a product of a polymerase chain reaction amplification. The detected nucleic acid may be utilised for diagnosis of disease, sequencing of nucleic acids, cutilised for diagnosis of disease, sequencing of nucleic acids is based on observing a colour change with the naked eye and is cheap, fast, simple, and robust, requiring no specialised or expensive equipment. The current sequence is that of the oligonucleotide which is related to a thiolomodified oligonucleotide-gold colloid conjugate probe of the invention.
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Best Local S
   29-JUL-1996
21-JUL-1997
29-JAN-1999
25-JUN-1999
13-JAN-2000
26-MAR-2000
26-APR-2000
26-JUN-2000
26-JUN-2000
11-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Park
                                                                                                                                                                                                                                                                                                                                            08-OCT-2002; 2002US-00266983
                                                                                                                                                                                                                                                                                                                                                                                                        06-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2003207296-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nanoparticle; gold; disease; forensic; paternity testing;
cell line authentication; gene therapy; ss; gold colloid conjugate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic thiol-modified oligo-gold colloid conjugate probe -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADI32905;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADI32905 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 24; SEQ ID NO 70; 206pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1520 АЛАЛЛАЛАЛАЛАГТАЛАЛ 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 AAAAAAAAAAAAAAAAA 18
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16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
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                               ; 2000US-0176409P.
; 2000US-0192699P.
; 2000US-0200161P.
; 2000US-00603830.
; 2000US-0213906P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
         2000US-0224631P
                                                                                                                                                                                         99US-00240755.
99US-00344667.
                                                                                                                                                                                                                                                    96US-0031809P.
97WO-US012783.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1; Length 20; Pred. No. 4.4e+02; O; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ç
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RESULT 781 ADK70840

ADK70840 standard;

DNA;

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1 ААДААЛАЛАЛАЛАЛАЛА 18

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TXAXAX EXEX EXXXXX

06-MAY-2004

(first entry)

ADK70840;

DNA preparation; 5' mRNA; linker synthesis; gene regulation; gene expression; ss; tag.

primer synthesis;

WO2003106672-A2 Unidentified 5' mRNA DNA preparation method related tag DNA sequence #8.

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S
                                                                                                                                                      CC having at least two types of nanoparticles, such as gold, having attached coligonucleotides and observing a detectable change brought about by CC Oligonucleotides and observing a detectable change brought about by CC hybridisation of the oligonucleotides on the nanoparticles with the CC nucleic acid. The method of the invention may be useful for detecting a CC nucleic acid, preferably viral RNA or DNA, bacterial DNA, a gene CC associated with a disease, a fungal DNA, barthetic DNA or RNA, a CC polymerase chain reaction amplification. The detected nucleic acid may be CC utilised for disgnosis of disease, sequencing of nucleic acid may be CC polymerase chain reaction amplification. The detected nucleic acid may be CC gene therapy. The method for detecting the nucleic acids is based on CC observing a colour change with the naked eye and is cheap, fast, simple, and requiring no specialised or expensive equipment. The current sequence is that of the synthetic thiol-modified oligonucleotide-gold CC colloid conjugate probe of the invention which is linked via a thiol
                                                    Query Match
Best Local
                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-JAN-2001; 2001US-00760500.
28-MAR-2001; 2001US-00820279.
09-APR-2001; 2001US-0282640P.
10-AUG-2001; 2001US-0027777.
09-OCT-2001; 2001US-0327864P.
07-DEC-2001; 2001US-00008978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-DEC-2000;
08-DEC-2000;
11-DEC-2000;
11-DEC-2000;
11-DEC-2000;
12-MAR-2001;
28-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detection of nucleic acid, e.g. viral RNA or DNA, comprises contacting nucleic acid with different types of nanoparticles having attached oligonucleotides and observing detectable change brought about by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Park
                                                                                                          Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (PARK/)
(TATO/)
                                                                                                                                           group to a gold nanoparticle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel method for detecting a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 18; SEQ ID NO 55; 206pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hybridization.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-059754/06.
   1520 AAAAAAAAAAAGTAAAA 1537
                                     16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MIRKIN C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PARK S.
TATON T A.
                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Taton TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-0254392P.
2000US-0254418P.
2000US-0255235P.
2000US-0255236P.
                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mirkin
                                                  1.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       S
                                   0
                                                                        Score 14.8;
                                                       Pred. No.
                                       Mismatches
                                                     4.4e+02;
                                                                      DB 1; Length 20
                                       2
                                       0
                                       Gaps
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RESULT 782
ADK69880/c
ID ADK698
XX ADK698
XX O6-MAY
XX Phosph
XX PH Key
FT modifi
FT
FT
XX US2003
XX US2003
XX US2003
XX IJ-JAL
PR 11-JAL
PR 11-JAL
PR 10-JAL
XX PA (COLE/
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              corresponding to a nucleotide sequence of a 5' end of an mRNA. The method is useful for synthesising a nucleotide sequence to be used as a linker or primer and selectively collecting multiple nucleic acid fragments containing information on the nucleotide sequences at the 5' end of multiple mRNA in a sample. The method is also useful for identifying regions in the genome, which are required for gene regulation and gene
                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                               Sulphurised oligonucleotide #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Preparing DNA fragment corresponding region of mRNA, by preparing nucleic sequence of 5' end of mRNA, cleaving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RIKE )
                             11-JAN-2000; 2000US-00481486.
10-JAN-2001; 2001WO-US000715.
                                                                                                   13-NOV-2003.
                                                                                                                               US2003212267-A1.
                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                      Phosphorothioate backbone; sulphurised oligonucleotide; ss
                                                                                                                                                                                                                                                                                                                                               06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                           ADK69880;
                                                                                                                                                                                                                                                                                                                                                                                                       ADK69880 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 0 A; 1 C; 10 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention comprises a method for preparing a DNA fragment corresponding to a nucleotide sequence of a 5' end of an mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 5; SEQ ID NO 40; 121pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hayashizaki Y, Carninci P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-JUN-2002; 2002JP-00171851.
12-AUG-2002; 2002JP-00235294.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-DEC-2003
                                                                       12-DEC-2002; 2002US-00181200
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  (COLE/) COLE D L.
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nee 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      545 rerecerecerecere 562
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rerererererecere 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present DNA sequence was used in an example of the
                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u></u>
                                                                                                                                                                                                /*tag=
                                                                                                                                                                      /mod_base= OTHER
/note= "Phosphorothioate backbone; 2'-0-methoxyethyl
                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                           residues"
                                                                                                                                                                                                                    . 20
                                                                                                                                                                                                                                                                                                                                                                                                     DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Harbers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          to nucleotide sequence of 5' end acid corresponding to nucleotide nucleic acid with restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.4e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
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ADK69885/c
ID ADK698
XX ADK698
AC ADK698
XX Sulphu
XX Phosph
XX Uniden
XX Uniden
XX Uniden
XX US2003
XX US2003
XX I1-JAN
PF 12-DE(
XX I1-JAN
PF 10-JAN
PF 10
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              phosphite intermediate; and oxidising the phosphite intermediate with an acetyl disulfide in an acetonitrile for a time to effect conversion of the phosphite intermediate to phosphorothioate. The invented method achieves high yields and greater efficiency. The present sequence is sulphurised oligonucleotide used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Preparation of phosphorothioate oligonucleotides involves oxidizing phosphite intermediate with acetyl disulfide in acetonitrile for time effect conversion of phosphite intermediate to phosphorothioate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to phosphorothioate oligonucleotides having nucleoside with 240 modification are prepared by phosphitylating 5'-hydroxyl of a nucleic acid moiety having a nucleoside with 2' modification in an acetonitrile containing solvent mixture to form a modification in an acetonitrile containing solvent mixture to form a
                                                      WPI; 2004-069376/07.
                                                                                                                                                                                                                                                                  11-JAN-2000;
10-JAN-2001;
                                                                                                                                                                                                                                                                                                                                           12-DEC-2002; 2002US-00181200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Phosphorothicate backbone; sulphurised oligonuclectide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sulphurised oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADK69885;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 12; SEQ ID NO 10; 8pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cole DL,
                                                                                                          Cole DL,
                                                                                                                                                        (COLE/) COLE D L.
(RAVI/) RAVIKUMAR V T.
(CHER/) CHERUVALLATH Z
                                                                                                                                                                                                                                                                                                                                                                                               13-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                US2003212267-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADK69885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RAVI/) RAVIKUMAR V T. (CHER/) CHERUVALLATH 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1520 АААААААААААСТАААА 1537
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAAAAAAAAAAAAAAAA
                                                                                                       Ravikumar VT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ravikumar VT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                               2000US-00481486
2001WO-US000715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "Phosphorothioate backbone; 2'-O-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%;
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                                                                                                          Cheruvallath
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 4.40
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1;
Pred. No. 4.4e+02;
                                                                                                          SZ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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Preparation of phosphorothicate oligonuclectides involves oxidizing

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RESULT 784
ADK74647/c
ID ADK746
XX ADK746
XX ADK746
XX Chimex
XX Chimex
XX Wavl.3
KW diabet
KW infant
XX WO2004
XX WO2004
XX HO2004
XX Chim
XX New ar
PT Navl.3
PT With h
PT disorc
XX Claim
XX Claim
XX The pa
CC The pa
CC The pa
CC special
CC special
CC neurog
CC headac
CC headac
CC headac
CC headac
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Best Local
                 The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, triggminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including headache; seizure disorder, including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to phosphorothioate oligonucleotides having nucleoside with 240 modification are prepared by phosphitylating 5'-hydroxyl of a nucleic acid moiety having a nucleoside with 2' modification in an acetonitrile containing solvent mixture to form a phosphite intermediate; and oxidising the phosphite intermediate with an acetyl disulfide in an acetonitrile for a time to effect conversion of the phosphite intermediate to phosphorothioate. The invented method achieves high yields and greater efficiency. The present sequence is sulphurised oligonucleotide used in the exemplification of the invention.
                                                                                                                                                                                                                                                                      New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-203785/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-AUG-2002; 2002US-0403416P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004016754-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Navl.3; Analgesic; Nootropic; Neuroprotective; diabetic neuropathy; arthritic pain; migraine l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric phosphorothioate oligonucleotide to target Nav1.3 #1981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADK74647 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     phosphite intermediate with acetyl disulfide in acetonitrile for effect conversion of phosphite intermediate to phosphorothicate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                               4; SEQ ID NO 1981; 417pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                        SL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 0 A;
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    neonatal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 C; 0 G; 20 T; 0 U; 0 Other;
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Pred. No. 4.4e+02;
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                                                                                                                                                                                                                               English
  epilepsy; or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; post-herpetic neuralgia;
headache;
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Query Match Best Local S Matches 16

Similarity

1.1%;

Score 14.8; DB 1 Pred. No. 4.4e+02 0; Mismatches

DB 1; Length 20;

<u>,</u>

Gaps

0

Sequence 20

BP;

0 A

1 C; 0 G; 19 T; 0 U; 0

Other;

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RESULT 785
ADK74188/c
ID ADK741
XX ADK741
XX ADK741
XX ADK741
XX AV1.3
CC CAMAY
XX Mav1.3
XW diabet
XX WO2004
XX WO2004
XX WO2004
XX I14-AUC
XX I4-AUC
XX I4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        á
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Best Local S
Matches 16
                                                                                      The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2 MOS wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence represents a chimeric phosphorothicate ol
2'MOE wings and a deoxy gap. Used during the antis
human Nav1.3 expression, the oligonuclectides are
different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nav1.3; Analgesic; Nootropic; Neuroprotective; diabetic neuropathy; arthritic pain; migraine h
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chimeric phosphorothicate oligonucleotide to target Navl.3 #1522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 1522; 417pp; English.
                                                               regions of the human Nav1.3 RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  epilepsy; ataxia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
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Pred. No. 4
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RESULT 786
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XX ADK749
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XX AV1.3
KW Giabet
KW infant
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RESULT 787
ADK75847
ID ADK758
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                                ADK75847 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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RESULT 788
ADK79524/c
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                            Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache;
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epilepsy; ataxia;
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Pred. No. 4.4e+02;
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                   14-AUG-2002; 2002US-0403416P
                                              14-AUG-2003; 2003WO-US025465
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                                                                                                                                           Synthetic
                                                                                                                                                                                       Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic diabetic neuropathy; arthritic pain; migraine headache;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 BP; 11 A; 1 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                             TTTTGATTTTTAATCACA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                 DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
                                                                                                                                                                                                                                                                                                                                 ВP
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length
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                                                                                                                                                                                                       neuralgia;
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RESULT 790
ADK74889/c
ID ADK748
XX ADK748
XX ADK748
XX Chimex
XX Chimex
XX Nav1.3
KW diabet
KW infant
XX Synthe
XX WO2004
XX FI 14-AU(
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XX I14-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, chapter in the pain from burns, migraine headache, cluster headache, mild-to-moderate pain from burns, migraine headache, cluster headache, mild-to-moderate pain neuropathy such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2°MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target odifferent regions of the human Navl.3 RNA.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                               14-AUG-2003; 2003WO-US025465.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric phosphorothicate oligonuclectide to target Nav1.3 #2223.
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                                                                                                                                                                                                                                                                                                (PHAA)
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                                                                                                                                                                                                                                                                                                   PHARMACIA CORP
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Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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Claim

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2223; 417pp; English.

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RESULT 791
ADK80788/c
ID ADK807
XX ADK807
XX ADK807
XX ADK807
XX ADK807
XX Chimer
XX Nav1.3
KW Gilabet
KW Infant
XX WO2004
XX WO2004
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
  The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navi.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navi.3. The compound and composition are useful for treating a disease or condition associated with Navi.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pair diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain,
                                                                                                                                                                                                                                                                                         New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav13, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
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nilarity 88.9%;
Conservative
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Pred. No. 4.4e
0; Mismatches
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ID MX74838/c
ID ADK748
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XX ADK748
XX ADK748
XX ADK748
XX Chimer
XX VAI Adabet
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Best Local
                                                       The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache, seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present seguence represents a chimeric phosphorothioate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target different regions of the human Navl.3 RNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-AUG-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nav1.3; Analgesic; Nootropic; Neuroprotective; diabetic neuropathy; arthritic pain; migraine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric phosphorothioate oligonucleotide to target Nav1.3 #2172.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2004016754-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1252
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16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TTTTGTTTTTTAATCAGA 1269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 2172; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2003WO-US025465
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Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; post-herpetic neuralgia;
headache;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
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                                                                                                                   Matches
                                                                                                                                           Query Match
Best Local (
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Best Local :
                                                                                                                                                                                                                                                                                 The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache, seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Roberds SL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-AUG-2002; 2002US-0403416P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PHAA ) PHARMACIA CORP.
                                                         1165
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                                                                                                                   16;
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16; Conserv
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                                                      AGTATTTGTTGAAATAGT 1182
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ilarity 88.9%;
Conservative
                                                                                                                                                                                                                                 BP; 7 A; 2 C; 4 G; 7 T; 0 U; 0 Other;
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                                                                                                                                           88.98;
                                                                                                                                                                    1.1%;
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Pred. No. 4.
                                                                                                                                        Score 14.8;
Pred. No. 4
                                                                                                                      Miematches
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4.4e+02;
2;
                                                                                                                                           4.4e+02;
                                                                                                                                                                       DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 20;
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RESULT

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ADK80450/c
ID ADK80450 standard; DNA; 20
XX
AC ADK80450;
XX
Chimeric phosphorothioate
XX
Chimeric phosphorothioate
XX
Mavl.3; Analgesic; Nootrop
XX
Chimeric neuropathy; arthi
XX
Infantile epilepsy; ataxi;
OS
Synthetic.
XX
26-FEB-2004.
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14-AUG-2002; 2002US-04034
XX
PF
14-AUG-2002; 2002US-04034
XX
PF
14-AUG-2003; 2003WO-US025.
XX
PF
14-AUG-2003; 2003WO-US025.
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PF
14-AUG-2003; 2002US-04034
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14
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ID ADM6
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AC ADM6
XC ADM6
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The
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                                                                 Plant gene polymorphism marker related primer, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 7784; 417pp; English.
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                                                                                                                                     03-JUN-2004
                                                                                                                                                                                                      ADM69507;
                                                                                                                                                                                                                                                                    ADM69507 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 10 A; 1 C; 3 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1252 TTTTGTTTTTTAATCAGA 1269
                                                                                                                                                                                                                                                                                                                                                                                                                                         19 TTTTGATTTTTAATCACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16;
variation mapping; mutation mapping; plant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8; DB 1;
Pred. No. 4.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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RRESULT 796
ADM69506
ID 9506
AC ADM695
XX ADM695
XX O3-JUN
DE Plant
XX Primez
KW Primez
KW Gene p
XX Synthe
XX Synthe
PN JP2003
XX PD 14-OCT
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       utilizing a genetic polymorphism marker as follows: (a) genomic DNA is prepared from a plant homozygously having a mutation to be an object of the mapping; (b) A forward primer I containing a base corresponding to the gene polymorphic maker of one ecotype plant, a forward primer 2 containing a base corresponding to the genetic polymorphism of the other ecotype plant and a reverse primer 3 based on the base sequence common with both the ecotype plants are prepared; (c) two kinds of oligonucleotides emitting fluorescence of different colors when the genetic polymorphism marker is detected are prepared; (d) an end of the primers 1, 2 and 3 and the two kinds of the oligonucleotides; (e) the fluorescence intensity emitted from the resultant reactional product is detected and (f) the position on the genome of the mutation site is detected and from the resultant reactional product is detected and from the results of detection site is
                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 1.1%;
Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                            ADM69506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a primer set and method for mapping least the variation/mutation part of a plant gene using a gene polymorphism marker. A mutation site of the plant gene is mapped by utilizing a genetic polymorphism marker as follows: (a) genomic DNA is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 7; SEQ ID NO 386; 120pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A primer set and method useful for mapping at least the variation/mutation part of a plant gene using a gene polymorphism marker.
                   14-OCT-2003
                                                      JP2003289885-A
                                                                                                                       Primer; variation mapping; mutation mapping; plant; gene polymorphism marker; ss.
                                                                                                                                                                         Plant gene polymorphism marker related primer, SEQ ID 385
                                                                                                                                                                                                           03-JUN-2004
                                                                                                                                                                                                                                                                              ADM69506 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer, used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-FEB-2002; 2002JP-00025338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-JAN-2003; 2003JP-00024620
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene polymorphism marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIKA ) RIKAGAKU KENKYUSHO
(SAIM-) SAI MEDIA KK.
(MATS/) MATSUI M.
                                                                                                                                                                                                                                                                                                                                                                                                   1462 AGAAGGTGAACAATTTCA 1479
                                                                                                                                                                                                                                                                                                                                                                     -
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NAKAZAWA M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 BP; 6
                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; 3 C; 7
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                                                                                                                                                                                                                                                                                                                                                                     18
                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; D
Pred. No. 4.4e
O; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.4e+02;
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RESULT 797
ADL33726/c
ID ADL3377
XX ADL337
XX ADL337
DT 03-JUN
XX LNA ol
XX Detect
XW Detect
XW Detect
XX Synthe
XX Key
FH Key
FH modifi
FT modifi
FT modifi
FT T
FT T
XX X
PN W02004
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Loc
Matches
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(SAIM-)
(MATS/)
                                                                                                          modified_base
                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                  LNA oligomer
                                                                                                                                                                                                                                                                                                                                                                              ADL33726 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 7; SEQ ID NO 385; 120pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A primer set and method useful for mapping at least the variation/mutation part of a plant gene using a gene polymorphism marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-FEB-2002; 2002JP-00025338
                   WO2004020575-A2
                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                    Detection; isolation; locked nucleic acid; LNA; ss
                                                                                                                                                                                                                                                                                                                  03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                 ADL33726;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1462 AGAAGGTGAACAATTTCA 1479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention relates to a primer set and method for mapping
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SAI MEDIA KK.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                               /*tag= p
/mod_base= OTHER
                                                                                                                                                                                      Location/Qualifiers
                                               /mod_base= OTHER
/note= "Optionally biotinylated or 5′ AQ2-HEG3, where AQ
is anthraquinone and HEG is hexa-ethylene glycol"
                                                                                            /*tag=
                                                                                                                             notes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 3 C; 6 G; 4 T;
                                                                                                                                                                                                                                                                                                                  entry)
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                                                                                                                             "Optionally
                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                ВP
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8;
                                                                                                                             LNA nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 Other;
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RESULT 798
ADL59686
ID ADL596
XX ADL596
XX ADL596
XX O3-JUN
XX Human
XX Cytost
KW Gene t
KW Gene t
KW angiog
KW angiog
KW angiog
KW angiog
KW angiog
KW RESM-1
KW BESM-1
KW BESM-1
KW RESM-1
KW RESM-1
FT modifi
FT modifi
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a method (M1) for detecting and/or isolating a nucleic acid having a homopolymeric sequence or repetitive element or conserved nucleotide sequence. (M1) comprises treating a sample containing nucleic acid compounds with an locked nucleic acid (LNA) oligonucleotide (LO) to thereby detect and/or isolate a nucleic acid having the homopolymeric sequence or repetitive element or conserved nucleotide sequence. (M1) is useful for detecting and isolating nucleic acids released from a lysed complex biological mixture comprising nucleic acids. The present sequence is a LNA oligomer, used to illustrate the
                                                                                                                                                                                                                                                                                       cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific molecule—1; ESM-1; ESM-1 related disorder; diabetes; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; antisense technology; ss.
                                                                     modified_base
                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                   Human ESM-1
                                                                                                                                                                                                                                                                                                                                                                                                                  03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                ADL59686;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acid oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detecting and/or isolating nucleic acid molecule having homopolymeric sequence or repetitive element or conserved nucleotide sequence involuteating sample containing nucleic acid compounds with locked nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JUN-2002; 2002US-0390928P
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                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADL59686 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 22; Page 51; 104pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EXIQ-) EXIQON AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1520
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP;
                                                                                                                                                                                                                                                                                                                                                                                   antisense oligonucleotide seqid 1935.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2003WO-IB006354
                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jacobsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                /mod_base= OTHER
/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
/*tag= c
/mod_base= OTHER
/not== "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                     mod_base= OTHER
                                                                                   note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ą
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.1%;
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <del>,</del>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, isoshaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonucleotide that can be used to modulate expression of
                                                                                                                                                                                                                        cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific molecule-1; ESM-1; ESM-1; ESM-1 diabetes; cancer; ischaemia; reperfusion injuangiogenic disorder; immunological disorder; cardiovascular disorder; angiogenic disorder; immunological disorder; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
                                                   modified_base
                                                                                                                                                                                                                                                                                                  Human ESM-1 antisense oligonucleotide seqid 1952.
                                                                                                                                                                                                                                                                                                                                                               ADL59703;
                                                                                                                                                                                                                                                                                                                                                                                            ADL59703
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; SEQ ID NO 1935; 555pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-248358/23.
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                                                                                                                             modified_base
                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                          neurological disorder; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        endothelial specific molecule-1 (ESM-1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Weinstein EJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1520 АЛАЛАЛАЛАЛАГТАЛАЛ 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAAAAAAAAGCACAA
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                                                                                                                                                                                                                                                                                                                                  (first
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                                                                     residues
                                                                                                                                              Location/Qualifiers
                                                                     /note= "OTHER= phosphorothioate residues are 5-methylcytidines"
                                                                                                                  /*tag=
                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.1%;
Dase= OTHER = 2'-methoxyethyl (2'-MOE)
                                                                                                  base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                                                                             BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.8;
Pred. No. 4
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                                                                                    All cytidine
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ADL59724
ID ADL59724
XX ADL597
XX ADL597
XX O3-JUN
DT O3-JUN
DX Human
XX Cytost
KW Cytost
KW Gene t
KW BSM-1
KW angiog
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Best Local
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                                                                                                                                              cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific moleculer; ESM-1; ESM-1; ESM-1; diabetes; cancer; ischaemia; reperfusion inju angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 16 A; 2 C; 1 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                               03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes a new antisense compound, having comprising 8-30 bp targeted to a nucleic acid encoding en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               encoding endothelial specific molecule-1 (ESM-1), userul ror prep. composition for treating e.g., diabetes, cancer or cardiovascular
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                                                               modified_base
                                                                                                                                                                                                                                             Human ESM-1 antisense oligonucleotide segid 1973
                                                                                                                                                                                                                                                                                                                                             ADL59724 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; SEQ ID NO 1952; 555pp; English
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                                                                                                                sapiens.
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/mod_base= OTHER
/note= "OTHER= phosphorothioate
residues are 5-methylcytidines"
                                                                                Location/Qualifiers
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/mod_base= OTHER
                                                   '*tag= b
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                                                                                                                                                                                                                                                                                                                                               20 BP
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Pred. No. 4.
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                    backbone.
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                                                                                                                                                                               reperfusion injury;
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endothelial
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                    cytidine
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RESULT 801
ADM93653/c
ID ADM936
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XX ADM936
XX ADM936
XX Human
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Best Local
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                                                                                                          Human; NOVX; PCR; ss; congenital heart defect; cardiomyopathy; atherosclerosis; hypertension; pulmonary stenosis; scleroderma; adenocarcinoma; haemophilia; graft-versus-host disease; cancer; neurodegemerative disorder; Alzheimer's disease; Parkinson's di multiple sclerosis; diabetes; obesity; bronchial asthma; acquired immunodeficiency syndrome; AIDS; Crohn's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
                                                                                                                                                                                                                                                                                                                 Human NOVX PCR
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                                                                                        infectious disease;
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16. .20
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/note= "OTHER=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
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05-NOV-2001
06-NOV-2001
09-NOV-2001
15-NOV-2001
21-NOV-2001
21-NOV-2001
29-NOV-2001
29-NOV-2001
04-DEC-2001
05-FEB-2002
15-MAY-2002
15-MAY-2002
16-MAY-2002
16-MAY-2002
06-AUG-2002
07-AUG-2002
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(ALSO/)
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PADIGARU M.
PATTURAJAN M.
PENA C B A.
PEYMAN J A.
RASTELLI L.
RIEGER D K.
ROTHENBERG M E
SHENOY S G.
SHIMKETS R A.
SMITHSON G.
SMITHSON G.
SMITHSON G.
SPADERNA S K.
STARLING G.
SPATRLING G.
                                                                                                                                                                                                        MEZES P S.
MILLER C E.
MILLET I.
MISHRA V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BARON M.
BOLDOG F L.
BULGESS C E.
CASMAN S J.
CHAPOVAL A.
DHANABAL M.
EDINGER S R.
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ALVAREZ E.
ANDERSON D W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    GORMAN L.
GROSSE W M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ELLERMAN K.
ETTENBERG S.
GANGOLLI E A.
GERLACH V.
                                                                                                                                                                                                                                                                                                         MALYANKAR U M.
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                                                                                                                                                                                                                                                                                                                                                                         KEKUDA R.
KHRAMTSOV N V.
                                                                                                                                                                                                                                                                                                                                                                                                                          HACKETT C.
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2001US-0335610P.
2001US-0333152P.
2001US-033312P.
2001US-0333912P.
2001US-0334309P.
2002US-034807P.
2002US-0380968P.
2002US-0381043P.
2002US-039148P.
2002US-039148P.
2002US-039148P.
2002US-0401676P.
2002US-0401676P.
2002US-0401676P.
2002US-0401676P.
2002US-0401693P.
2002US-0401693P.
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2001US-0333072P.
2001US-0345398P.
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RESULT 802
ADM13992/c
ID ADM13992/c
XX
ADM139
XX
AC ADM139
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DE Human
XX
Chimer
XW micros
XW micros
XW micros
XW immuno
XW immuno
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                                                                                                                                                                                                                                                                                                                                                                                              8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to human NOVX polypeptides and the polynucleotides concoding them. The NOVX polypeptides and polynucleotides are useful for determining the presence of or predisposition to a disease associated with altered levels of the sequences in a mammalian subject, and for treating or preventing a pathology associated with NOVX. The polypeptides, polynucleotides and antibodies that bind immunospecifically to the polypeptides are useful for treating or preventing disorders or syndromes such as congenital heart defects, cardiomyopathy, atherosclerosis, hypertension, pulmonary stenosis, scleroderma, cadenocarcinoma, haemophilia, graft-versus-host disease, cancer, neurodegenerative disorders, Alzheimer's disease, Parkinson's disease, multiple sclerosis, diabetes, obesity, bronchial asthma, acquired immunodeficiency syndrome (AIDS), Crohn's disease, Infectious disease, anorexia and immune disorders. This sequence represents a PCR primer used to amplify a human NOVX polynucleotide of the invention. Note: The sequence data for this patent; s also available from USPTO at
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 16
               chimeric; antisense oligonucleotide; phosphorothioate; human; microseomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microseomal prostaglandin E2 synthase; mPGES-1; protectic; antidiabetic, microseomal prostaglandin; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alsobrook JP, Alvarez E, Anderson DW, Baron M, Boldog FL;
Burgess CE, Casman SJ, Chapoval A, Dhanabal M, Edinger SR, Eisen
Ellerman K, Ettenberg S, Gangolli EA, Gerlach V, Gorman L;
Grosse WM, Guo X, Hackett C, Ji W, Kekuda R, Khramteov NV;
Lepley DM, Li L, Macdougall JR, Malyankar UM, Mazur A, Mcqueeney
Mezes PS, Miller CE, Millet I, Mishra V, Padigaru M, Patturajan
Pena CEA, Peyman JA, Rastelli I, Rieger DK, Rothenberg ME;
Shenoy SG, Shimkets RA, Smithson G, Spaderna SK, Starling G;
Spytek KA, Stone DJ, Tchernev VT, Twomlow N, Vernet CAM;
Zerhusen BD, Voss EZ, Zhong M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ZERH/)
(VOSS/)
(ZHON/)
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(TCHE/)
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                                                                                                                                                  Human mPGES-1 chimeric antisense oligonucleotide SEQ ID
                                                                                                                                                                                         01-JUL-2004
                                                                                                                                                                                                                              ADM13992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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TCHERNEV V T.
TWOMLOW N.
VERNET C A M.
ZERHUSEN B D.
VOSS E Z.
ZHONG M.
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                                                                                                                                                                                                                                                                        antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                         Sequence 20
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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to CC 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, cardiant, neuroprotective, cardiant, neuroprotective, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or
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/note= "phosphorothioate linkages and all cytidine
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                                                                                   New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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== "2'-O-methoxyethyls"
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Pred. No. 4.
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RESULT 805
ADM1408/c
ID ADM140
XX ADM140
XX ADM140
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Matches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; carddant; neuroprotective; antinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                  modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                     note= "phosphorothicate linkages"
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== "2'-O-methocyethyls"
                                                                                                                                                      base= OTHER
== "2'-O-methoxyethyls'
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No. 4
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08-APR-2004

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   RESULT 806
ADM14002/c
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Best Local (
                                                                                                                                           chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; cardiant E2 synthase; mhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9344.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, necroprotective.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                   Human mPGES-1
                                                                                                                                                                                                                                                                                                                                                     01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                              ADM14002 standard; DNA; 20 BP
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                                                                        Synthetic.
                    modified_base
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                                                                                                                                                                                                                                                                                                                 chimeric antisense oligonucleotide SEQ
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Location/Qualifiers
1. .20
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                                                                                                                            neurological disorder;
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Pred. No. 4.4e
0; Mismatches
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RESULT 807
ADM14090/c
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SEXEXEXE
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                                                                                                                                                                                                                                           Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                           Query Match
Best Local :
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               Human mPGES-1 chimeric antisense
                                            01-JUL-2004
                                                                          ADM14090
                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ischemia.
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                                                                                                      ADM14090
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                                                                                                      standard; DNA; 20
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                                                                                                                                                                                                                                             Conservative
                                             (first entry)
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/mod_base= OTHER
'note= "2'-O-methoxyethyls"
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residues are 5-methylcytidines"
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note= "2'-O-methocyethyls"
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                                                                                                                                                                                                                                                          88.9%;
                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                         Score 14.8;
                                                                                                                                                                                                                                                             Pred.
                                                                                                                                                                                                                                             Mismatches
              oligonucleotide
                                                                                                                                                                                                                                                           No.
                                                                                                                                                                                                                                                           4.4e+02;
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                 NO:277
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The present sequence represents a chimeric antisense oligonucleotide trargeted to human microsomal prostaglandin E2 synthase (mpCES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to comprise the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antistanse oligonucleotides and antisense compounds have cytostatic, antistanse oligonucleotides and antisense compounds have cytostatic, ophthalmological, immunomodulator, cardiant, neuroprotective, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
Query Match
Best Local S
Matches 16
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                       The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mBGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion, injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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Pred. No. 4.4e+02
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ADM14017/c
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothioate;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1
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                  modified_base
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                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                               cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human mPGES-1 chimeric antisense oligonucleotide SEQ ID
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
/*tag= c
                                                                                                               /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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Pred. No. 4.4e+02;
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RESULT 811
ADM14018/c
ID ADM140
XX ADM140
AC ADM140
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XX OLimer
KW Micros
KW Micros
KW immuno
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC human mpGES-1 gene is located on chromosome 9, more specifically to G2 434.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybriddise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, antistinflammatory, neuroprotective, nootropic, antistrhritic, vasotropic, CC ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense or CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 16
                   chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1 inhibitor; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1; mtdiabetic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                          Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:205
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence represents a chimeric antisense oligonucleotide human microsomal prostaglandin E2 synthase (mPGES-1). The
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Pred. No. 4.4e+02;
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                                                                                    Matches
                                                                                                     Query Match
Best Local
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modified_base
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/note= "phosphorothioate linkages and
residues are 5-methylcytidines"
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/note= "2'-O-methoxyethyls"
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RESULT 812 ADM14088/c ID ADM14088 standard; DNA;

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9d34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabletic, immunomodulator, cardiant, necoprotective,
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antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.qinflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 275; 132pp; English.
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note= "2'-O-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "phosphorothioate linkages and residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mod base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          base= OTHER
== "2'-O-methoxyethyls"
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SXX
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ADM14257/c
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1 imbibitor; microsomal prostaglandin E2 synthase imbibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; immunomodulator; cardiant; neuroprotective; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                      Claim 4; SEQ ID NO 444; 132pp; English
                                                                                                                                                                                                                   25-SEP-2003; 2003WO-US030374
                                                                                                                                                                                                                                              08-APR-2004.
                                                                                                                                                                                                                                                                      WO2004028458-A2
                                                                                                                                                                                                                                                                                                                                   modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:444
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ophthalmic, immunological, cardiovascular or neurological disorder
                                                                                                               WPI; 2004-305094/28.
                                                                                                                                                                 (PHAA ) PHARMACIA CORP
                                                                                                                                                                                         25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                       16. .20
                                                                                                                                                                                                                                                                                                                                                                                                     /note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                           /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                note= "2'-0-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                 mod_base= OTHER
                                                                                                                                                                                                                                                                                                note= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                             _base= OTHER
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Pred. No. 4.
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                                                                           e.g.,
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The present sequence represents a chimeric antisense oligonucleotide

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RESULT 814
ADM14000/c
ID ADM140
XX ADM140
XX ADM140
XX ADM140
XX Chimer
KW micros
KW m
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Best Local
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                                     25-SEP-2003; 2003WO-US030374.
                                                                                                   08-APR-2004.
                                                                                                                                                                    WO2004028458-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                        16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                   note= "2'-0-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                                                         mod_base= OTHER
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                                                                                                                                                                                                                                 _base= OTHER
== "2'-O-methoxyethyls"
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Pred. No. 4.4e+02;
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FTT FTT XXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to comprise the present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 by targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and comprise the sexpression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisintlammatory, neuroprotective, noutropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, noutropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, noncomposition that activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or repertusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 16
                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                  modified_base
                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                    Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:193
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM14006 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 187;
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                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1520 AAAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                Location/Qualifiers
1. .20
/*tag= b
/note= "phosphorothioate linkages and all cytidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88.9%;
                  base= OTHER
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Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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RESULT 816
ADM14014/c
ID ADM140
XX
AC ADM140
XX
DT 01-JUL
XX
XX
DB Human
XX
KW chimer
KW micros
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                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to CC 9934.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mPGES-1, which specifically hybridise with the nucleic acid encoding CC inhibits its expression; (2) a method of inhibiting the expression of MPGES-1 in cells or tissues, and (3) a method of treating an animal CC having a disease or condition associated with mPGES-1. MPGES-1 chimeric CC antistagence or condition and antisense compounds have cytostatic, CC antistinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, CC antistalmanatory, neuroprotective, nootropic, antistarthritic, vasotropic, CC ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's CC ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                    Query Match
chimeric; antisense oligonucleotide; phosphorothioate; human;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
                                                  Human mPGES-1
                                                                                   01-JUL-2004
                                                                                                                    ADM14014
                                                                                                                                                     ADM14014 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                                                                                     BP; 0 A; 0 C; 0 G; 20 T; 0 U;
                                                                                                                                                                                                                                                                                                       Conservative
                                                                                   (first entry)
                                                  chimeric antisense oligonucleotide SEQ ID
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/note= "2'-O-methoxyethyls"
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Pred. No. 4.4e+02;
0; Mismatches 2
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                                                                                                                                                                                                                                                                                                                                  DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                     0 Other;
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1520 AAAAAAAAAAAAGTAAAA 1537

Matches Query Match Best Local :

16;

Conservative

0; Mismatches Pred. Score 14.8; ŏ.

4.4e+02 DB 1;

Length

20

Indels

0

Gaps

0

Similarity

88.9%;

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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The Chuman mpGES-1 gene is located on chromosome 9, more specifically to CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC mpGES-1 in cells or condition associated with mpGES-1. chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, or antiantial method of the composition con be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's content of the condition associated with mpGES-1 e.g., inflammation, Alzheimer's content of the condition associated with mpGES-1 e.g., inflammation, Alzheimer's content of the condition associated with mpGES-1 e.g., inflammation, Alzheimer's content of the condition associated with mpGES-1 e.g., inflammation, Alzheimer's content of the condition of the conditio
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; SEQ ID NO 201; 132pp; English.
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                                                                                                    ophthalmic,
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     B₽;
                                                                                              immunological, cardiovascular or
0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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residues are 5-methylcytidines"
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                                                                                              neurological
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           targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibite its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric
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                                                                                                                                                                                                                       New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.c inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
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                                                                                                                                           The present sequence represents a chimeric antisense oligonucleotide
                                                                                                                                                                           Claim 4; SEQ ID NO 207; 132pp; English.
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "2'-O-methoxyethyls"
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antisense oligonucleotides and antisense compounds

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New antisense compound, having a sequence targeted

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                          Key
modified_base
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                                                     Gierse JK;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:178.
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                                                                                                           25-SEP-2002; 2002US-0413549P.
                                                                                                                                       25-SEP-2003; 2003WO-US030374
                                                                                                                                                                    08-APR-2004.
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                              /mod
                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                  . 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.1%;
                                                                                                                                                                                                                        base= OTHER
"2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                              base= OTHER
                                                                                                                                                                                                                                                                                "2'-O-methocyethyls"
                                                                                                                                                                                                                                                     a
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Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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RESULT 819
ADM14003/c
ID ADM140
AC ADM140
AC ADM140
AC ADM140
AC ADM140
AC ADM160
AC A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CI inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal compounds and condition associated with mpGES-1 chimeric cantisense oligonuclectides and antisense compounds have cytostatic, CC antisense oligonuclective, nootropic, antianthritic, vasotropic, antiinflammatory, neuroprotective, nootropic, antianthritic, vasotropic, cophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound C can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 e.g., inflammation, Alzhaimer's C condition associated with mpGES-1 e.g., inflammation, Alzhaimer's compound can be used as rthritis, diabetes, cancer, ischaemia or reperfusion injury, or cophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzhelmer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:190.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADM14003 standard; DNA; 20
                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                     modified_base
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                                                                            /mod_base= OTHER
/note= "2'-O-methocyethyls"
16. .20
                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "phosphorothicate linkages and
reaidues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
/mod_base= OTHER
/note= "2'-O-methoxyethyls"
                                                            /*tag= c
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Pred. No. 4.
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E2 synthase (mPGES-1). Th
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpdES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to CC 934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, contininflammatory, neuroprotective, nootropic, antistification, immunomodulator, cardiant, neuroprotective, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.d inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-APR-2004.
Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; SEQ ID NO 190; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-SEP-2002; 2002US-0413549P
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밁
                                      Matches
                                                         Query Match
                                                Best Local
           1520 AAAAAAAAAAAAGTAAAA 1537
 20 AAAAAAAAAAAAAAAAAAA
                                       16;
                                                Similarity
                                      Conservative
                                               88.9%;
                                                         1.18;
                                      0; Mismatches
                                                         Score 14.8;
                                                Pred. No. 4.4e+02
                                                           DB 1;
                                                         Length
                                      <u>;</u>
                                       Gaps
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RESULT 820
ADM14005/c
ID ADM140
XX
AC ADM140
XX
AC ADM140
XX
DT 01-UUL
XX
Human
XX
Chimer
KW micros
KW micros
KW immuno
                                                                                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcBS-1; mpcBS-1 inhibitor; microsomal prostaglandin E2 synthase; mpcBS-1; mcgBS-1 inhibitor; cardiant E2 synthase inhibitor; cytostatic; antidiabetic immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                            Human mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                     01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADM14005;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADM14005 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                            chimeric antisense oligonucleotide SEQ ID
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NO:192

antidiabetic;

cardiovascular

disorder; neurological disorder;

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RESULT 821
ADM14246/c
ID ADM142
XX
AC ADM142
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Best Local :
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   ADM14246;
                                   ADM14246 standard; DNA; 20
                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 192; 132pp; English.
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                                                                                                                                                                                                 Local Similarity
                                                                                                                                                  1520
                                                                                                                                                                                  16;
                                                                                                                                                                                                                                                20
                                                                                                                                                  AAAAAAAAAAAGTAAAA 1537
                                                                                                                   AAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                                BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  *tag=
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:esidues are S-methylcytidines"
                                                                                                                                                                                                 1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 _base= OTHER
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== "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                base= OTHER
                                                                                                                                                                                <u>.</u>
                                                                                                                                                                                               Score 14.8; DB 1
Pred. No. 4.4e+02
                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                              Length 20
                                                                                                                                                                                   Indels
                                                                                                                                                                                0,
                                                                                                                                                                                  Gaps
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antidiabetic, immunomodulator, cardiant, neuroprotective, vasotropic, antiarthritic, vasotropic, antiarthritic, vasotropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation, Alzheimer's mediasese, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or

immunological, cardiovascular or neurological disorder

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targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9324.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidishetic immunromodalists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                   The present sequence represent targeted to human microsomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gierse
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                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 433; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-305094/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-SEP-2003; 2003WO-US030374.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2004028458-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ischemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
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                                                                                                                                                                                                                                                                                                                                       sequence represents a chimeric antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         base= OTHER
"2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "2'-0-methocyethyla"
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RESULT 822
ADM1395/c
IDM 1395/c
IDM 1395/c
ADM13995/c
ADM139D
ADM139D
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzhelmer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to
                                                                                                                                                                                     New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                          WPI; 2004-305094/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-SEP-2002; 2002US-0413549P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUL-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADM13995 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                (PHAA ) PHARMACIA CORP.
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                                                                                                              SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
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llarity 88.9%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              B₽;
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/note= "2'-O-methocyethyls"
16. .20
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                                                                                                         182; 132pp; English.
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"2'-O-methoxyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T; 0 U; 0 Other;
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                                                                                                                                                                                                                       e.g.,
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RESULT 823
ADM14011/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, antiminal antiverse compounds acradiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                  chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; inmunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                             WO2004028458-A2
                                                                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADM14011;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                  /mod_base= OTHER
/note= "phosphorothicate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                        /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                         note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                         base= OTHER
== "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.9%;
                                                                                                                                                                                                           "2'-O-methocyethyls"
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0; Mismatches
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    antisense compounds,

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25-SEP-2002; 2002US-0413549P 25-SEP-2003; 2003WO-US030374.

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RESULT 824
ADM14240/c
ID ADM142
XX ADM142
XX O1-JUL
XX Chimer
KW Chimer
KW Micros
KW Micros
KW Micros
KW Immuno
KW I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9q34.3. The present invention also describes: (1) shipperson, or condition and observables: (1) shipperson, which specifically hybridise with the nucleic acid encoding CC mpoES-1, which specifically hybridise with the nucleic acid encoding CC inhibits its expression; (2) a method of inhibiting the expression of CC mpoES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpoES-1. MpoES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, and carthritic, vasotropic, CC antinflammatory, neuroprotective, nootropic, antistribritic, vasotropic, cophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpoES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpoES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPCES-1; mPCES-1 inhibitor; microsomal prostaglandin E2 synthase; mPCES-1; mPCES-1 inhibitor; cardiant; neuroprotective; antinflammatory; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
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                                                                                                                                                                                                                                                                                                                                                                                                                         immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
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                                                                                                                                                                                                                                                                                                                                                                                            disorder; neurological disorder; ss.
                                              residues are
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                                           /note= "phosphorothioate linkages and
residues are 5-methylcytidines"
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RESULT 825
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents a chimeric antisense oligonucleotide cargeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to compare the present invention also describes: (1) antisense compounds, and in graph of the present invention also describes: (1) antisense compounds, mpGES-1, which specifically hybridise with the nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding carbibite its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues, and (3) a method of treating an animal carbing a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 e.g., inflammation, Alzheimer's carbing a cartifical displace, arthritis, diabetes, cancer, isbaemia or reperfusion injury, or
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Best Local :
chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin £2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin £2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;
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                                                                                      Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:196
                                                                                                                                                          ADM14009;
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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CC The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC maying a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, contininflammatory, neuroprotective, nouroprotective, antisinflammatory, neuroprotective, nouroprotective, cCC antidiabetic, immunomodulator, cardiant, neuroprotective, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 196; 132pp; English.
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic imminomedalistor.
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                                                                                                                                                                                                                                         ischemia
                                                                                                                                                                                                                                                                                                                                                                                (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                    2004-305094/28.
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                 immunomodulator,
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16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note= "2'-O-methoxyethyls"
 neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry.
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                 cardiant, neuroprotective,
    antiarthritic,
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RESULT 827
ADM14089/c
ID ADM140
XX ADM140
XX ADM140
XX Chimer
Chimer
KW Chimer
KW Micros
KW Micros
KW Immuno
KW Immu
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; mtcrosomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.c inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                            WPI; 2004-305094/28.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUL-2004
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                                                                                                                                                                                                                                                (PHAA ) PHARMACIA CORP
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/note= "2'-O-methoxyethyls"
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Pred. No. 4.4e+02;
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RESULT 828
ADM14016/c
ID ADM140
XX ADM140
XX ADM140
XX Chimer
KW Chimer
KW Micros
KW Micros
KW Immuno
KW I
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Best Local S
Matches 16
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                    /*tag= c
/mod_base= OTHER
/note= "2'-O-methoxyethyls"
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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.e= "2'-O-methocyethyls"
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Pred. No. 4.4e+02
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ARESULT 829
ADM14075/c
ID ADM140
XX ADM140
XX ADM140
DT 01-JUL
XX Human
XX Chiner
KW micros
KW micros
KW micros
KW immuno
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Best Local &
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                                                                                                  chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                        Homo sapiens
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RESULT 830
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AC ADM141
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OT 01-JUL

standard; DNA;

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ADM14189; ADM14189

01-JUL-2004

(first entry)

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                                                                                                                                                    The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The Chiman mpGES-1 gene is located on chromosome 9, more specifically to CC 9q34.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, CC antisense oligonucleotides and antisense compounds have cytostatic, contininflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's contribution associated with mpGES-1 e.g., inflammation, Alzheimer's contribution injury, or condition associated with mpGES-1 e.g., inflammation, Alzheimer's contribution injury, or conditions are composition for treating and cardiovascular activities.
                                                                           Matches
                                                                                           Query Match
Best Local
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                                                                                                                                                    Sequence 20
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/note= "2'-O-methocyethyls"
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Query Match Best Local S Matches 16

Similarity

1.1%;

Score 14.8; DB 1; Pred. No. 4.4e+02; Mismatches

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9344.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal cannot be used as a method of antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, cardiant, neuroprotective, or ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischemia or reperfusion injury, or other propersion of the condition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
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16. .20
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e= "2'-O-methocyethyls"
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0 Other

immunological,

cardiovascular

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                                                                                                                                                                             Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chimeric; antisense oligonucleotide; phosphorothioate; human;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encodi
                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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residues are 5-methylcytidines"
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RESULT 832
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      chimeric; antisense oligonucleotide; phosphorothioate; human; mkcrosomal prostaglandin E2 synthase; mpEES-1; mpEES-1 inhibitor; microsomal prostaglandin E2 synthase; mpEES-1; mpEES-1; mtidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                           modified_base
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                                  (PHAA ) PHARMACIA CORP
                                                                                               25-SEP-2002; 2002US-0413549P.
                                                                                                                                                        25-SEP-2003; 2003WO-US030374.
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llarity 88.9%;
Conservative
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residues are 5-methylcytidines"
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                                                                                                                                                                                             modified_base
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Similarity 88.9%;
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Pred. No. 4.4e+02;
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                             chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory;
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                 neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
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therapy; inflammation;
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Query Match
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Matches 16
                                                                                                                                                                                                                                          antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiinflammatory, neuroprotective, nootropic, antianthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperiusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric
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                                                                                                                                                                                                                      ophthalmic,
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                                                                                                                                                                                                                                                   (PHAA)
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                                                                                                                                                                                                                                                         PHARMACIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; DNA;
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residues are 5-methylcytidines"
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ss, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC human mpGES-1 gene is located on chromosome 9, more specifically to CC 934.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antisinflammatory, neuroprotective, nootropic, antistinflammatory, neuroprotective, nootropic, antistinflammatory. Immunomodulator, and cardiovascular activities, and can CC ophthalmological, immunomodulatory and cardiovascular activities, and can CC can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC conthibute; diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic immunofical activities and can composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC conthibute; diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic immunofical activities and cancer, ischaemia or reperfusion injury, or ophthalmic immunofical activities and cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:208
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                                                                                                                                      16. .20
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residues are 5-methylcytidines"
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/note= "2'-O-methoxyethyls"
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to Q934.3. The present invention also describes: (1) antisense compounds, CC mPGES-1, which specifically hybridise with the nucleic acid encoding CC inhibits its expression; (2) a method of inhibiting the expression of CC mPGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mPGES-1. MPGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound be used for preparing a composition for treating a disease or CC condition associated with mPGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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Best Local S
Matches 16
                                                                                                                                                                                                                                       chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1; mpGES-1; mpGES-1; mpGES-1; mpGES-1; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaematic propertusion injury; ophthalmic disorder; immunological disorder;
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                                                                                                                                             Homo sapiens
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Location/Qualifiers
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Best Local
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Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:200.
                             01-JUL-2004
                                                                                                                                                                                                                                                                                    Sequence
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to g934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding conhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisinse oligonucleotides and antisense compounds have cytostatic, and cantisinflammatory, neuroprotective, nootropic, antistrict, vasotropic, antistrictal, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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Query Match Best Local Similarity

1.1**%**; 88.9**%**;

Score 14.8; DB 1; Pred. No. 4.4e+02;

DB 1;

Length

20;

Sequence 20

BP; 0 A; 0 C; 0 G;

20 T; 0 U; 0 Other;

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RESULT 840
ADM14019/c
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           The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and
                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                          Claim 4; SEQ ID NO 206; 132pp; English.
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residues are 5-methylcytidines"
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Matches 16
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                                Gierse JK;
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es are 5-methylcytidines"
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Pred. No. 4.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzhelmer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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e= "2'-O-methocyethyls"
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Mismatches 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin 82 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to CC 934.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC maying a disease or condition associated with mpGES-1 wpGES-1 chimeric CC antisense oligonuclectides and antisense compounds have cytostatic, cantidiabetic, immunomodulator, cardiant, neuroprotective, antistinflammatory, neuroprotective, nootropic, antistribities, and can CC conthibation associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's conthibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's conthibitors, diabetes, cancer, ischaemia or reperfusion injury, or conthibitis, diabetes, cancer, ischaemia or reperfusion injury, or conthibitis, diabetes, cancer, ischaemia or reperfusion injury, or
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microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibit microsomal prostaglandin E2 synthase inhibitor; cytostatic; ar immunomodulator; cardiant; neuroprotective; antinfilammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthal immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder
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                                                                                                                                                                                                Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:180
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                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate;
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/mod_base= OTHER
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                                                                                                                                                                                                                              antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                    1.1%;
                                                                                                                                                                                    0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                 cardiovascular or neurological
                                                                                                                        0;
                                                                                                                                        Score 14.8; DB 1;
Pred. No. 4.4e+02;
                                                                                                                         Mismatches
                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin EZ synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 by targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antistisense oligonucleotides and antisense compounds have cytostatic, antistinflammatory, neuroprotective, nootropic, antistifical, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound cardiovascular activities, and can be used for preparing a composition for treating a disease or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antidiflammatcory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory, cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic avenceding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 185; 132pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:185
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 can be used condition as
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                                                                                                                                                                                                                                                                                                                               ischemia.
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 ed for preparing associated with
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residues are 5-methylcytidines"
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== "2'-O-methocyethyls"
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a composition mPGES-1 e.g.,
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RESULT 845
ADM14007/c
ID ADM140
XX ADM140
XX ADM140
XX Chimer
KW chimer
KW micros
KW m
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                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-2003; 2003WO-US030374.
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                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                               25-SEP-2002; 2002US-0413549P
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                          Claim 4; SEQ
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residues are 5-methylcytidines"
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Pred. No. 4.
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RESULT 846
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzhelmer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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modified_base
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                               /note= "
16. .20
                                                                                                                                                                                                                                                                                  /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                             /*tag= c
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                  /note = "phosphorothioate linkages residues are 5-methylcytidines"
                                                                                                                                                                                                                    _base= OTHEK
.e= "2'-O-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.1%;
                                                                                                                                                                                                                                                                                                                                                                                    _base= OTHER
                                                                                                                                         "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8; UL
. 4.4e+02;
2;
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                                                                                                                                                                                                                                                                                                                                                             a11
                                                                                                                                                                                                                                                                                                                                                           cytidine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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25-SEP-2003; 2003WO-US030374.

08-APR-2004.

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ADM14216/c
ID ADM142
XX ADM142
XX ADM142
XX ADM142
XX ADM16
DT 01-JUL
XX Chimer
XX Chimer
XX Immuno
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence represents a chimeric antisense oligonucleotide caracteristically to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to compare the present invention also describes: (1) antisense compounds, caracteristically hybridise with the nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding caracteristic its expression; (2) a method of inhibiting the expression of caracteristic its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal caracteristic immunomodulator, cardiant, method of treating an animal cantisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulator, and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischemia or reperfusion injury, or cardiant, arthritis, diabetes, cancer, ischemia or reperfusion injury, or cardiant in the cardiant or reperfusion injury, or cardiant in the cardiant or reperfusion injury, or cardiant in the cardiant 
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                       microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; microsomal prostaglandin E2 synthase intri-cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                       Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                      modified_base
                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothicate; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-JUL-2004
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                                                                                                                                                                                                                                                 sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunological, cardiovascular or neurological disorder
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                                                                                                                    Location/Qualifiers
                                     /*tag=
    /mod_base= OTHER
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Pred. No. 4.4e+02;
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ADO46424/c
ID ADO464
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AC ADO464
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AC ADO464
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C BT 15-JUL
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DE Human AX
XX
Human;
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                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compound, having a sequence targeted to a nucleic and encoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 4; SEQ ID NO 403; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gierse
                                  Human oligonucleotide #1790.
                                                                   15-JUL-2004
                                                                                                    ADO46424;
                                                                                                                                    ADO46424 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                       Sequence 20
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                                                                                                                                                                                                                                                      1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                         20 AAAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                                                                       16;
                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                      BP; 0
                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "phosphorothioate linkages and
residues are 5-methylcytidines"
1..5
                                                                                                                                                                                                                                                                                                                                                         A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                       1.1%;
                                                                                                                                     20
                                                                                                                                     ₽₽
                                                                                                                                                                                                                                                                                       <u>,</u>
                                                                                                                                                                                                                                                                                                         Score 14.8; DB 1.
Pred. No. 4.4e+02
                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                         Indels
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                                                                                                                                                                                                                                                                                       Gaps
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Human; ss;

interleukin-4 receptor; IL-4; interleukin-5 receptor;

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chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)

-5 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM,

C tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention

also relates to a method of screening a candidate compound that binds to

one or more nucleic acid target(s) or expressed product(s), for the

prevention and/or treatment of a respiratory or lung disease. The

coligonucleotides are useful for reducing or inhibiting expression of a

gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

CCR1, CCR3, Eotaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a,

cryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are

useful for preventing or treating a respiratory or lung disease. The

crespiratory or lung disease is associated with hyper-responsiveness to

and/or increased levels of, adenosine and/or levels of adenosine A

receptor(s), and/or asthma and/or lung allergies associated with

cinflammation or an inflammatory disease. The respiratory or lung disease

is chosen from airway inflammation, allergies associated with

cystic fibrosis (CF), chronic obstructive pulmonary disease (COPD),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PD84 A; PD84 B; PD84 C; PD84 D; respiratory disease; lung disease; hyper-responsivenese; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; alivway inflammation; allergy; impeded respiration, cystic fibrosis; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
Sequence 20
                                                                                                          allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shahabuddin
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(SAND/)
(TANG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-APR-2002;
23-APR-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (WILL/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SHAHABUDDIN S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MILLER S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGUILAR D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TANG
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in S, Lu H,
   BP; 0 A;
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2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ID NO 1791; 174pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cong
1 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                      This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tang
ong H;
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1520 AAAAAAAAAAAAGTAAAA 1537

Matches Query Match

16;

Conservative

0; Mismatches Score 14.8; DB 1; Pred. No. 4.4e+02;

Local Similarity

1.1%;

DB 1;

Length Indels

0

Gaps

0

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RESULT 849
AD003711
ID AD0037
XX
AD0037
XX
AC AD0037
XX
DT 29-JUL
XX
Raman
KW Raman
KW SERS;
XX
SYNTHE
PN US2004
XX
PR 07-MAY
PR 14-JUN
XX
PR 14-JUN
XX
PR (GAOY)
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                                                                                                                                                                                                                                                                                          The invention describes a reagent (I) comprising a particle bound to at CC least one Raman label and a specific binding member, where the Raman CC label can be activated to provide a surface-enhanced Raman scattering (SERS) effect or comprising a specific binding member having two or more different Raman labels bound to it. Also described are: a test kit (II), comprising (I) in one container and a silver, gold or copper Raman cCC (III), having a bundle of optical fibres terminating with ends of the optical fibre, where a several of the optical fibres have (I) located at the ends of the optical fibre. (I) is useful for: detecting for the presence or absence of one or more target analytes in a sample, the carget analytes having at least two binding sites; detecting the presence of the nucleic acid having at least two portions; and for screening one or more more daid having at least two portions; and for screening one or more specific receptors. This sequence represents an oligonucleotide cassociated with the SERS-based detection analyte detection method.
                                                                                                                          Matches
                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Reagent, useful for detecting target analyte e.g., nucleic acid, comprising particle having bound to at least one Raman label, which activated to provide surface-enhanced Raman scattering effect, specific binding member.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-MAY-2002; 2002US-0378538P.
28-MAY-2002; 2002US-0383630P.
14-JUN-2002; 2002US-00172428.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Raman label; specific binding member; surface-enhanced Raman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SERS-based analyte detection oligonucleotide segid 31.
                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; SEQ ID NO 31; 55pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-418413/39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MIRK/) MIRKIN C
(CAOY/) CAO Y.
(JINR/) JIN R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-MAY-2003; 2003US-00431341.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004.
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                                                           1520 AAAAAAAAAAAGTAAAA 1537
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                                                                                                                          16;
   1 AAAAAAAAAAAAAAAAAAAA
                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cao Y,
                                                                                                                                                                                                                                           BP;
                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                           20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                   1.1%;
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                                                                                                                       Score 14.8; DB 1;
Pred. No. 4.4e+02;
0; Mismatches
                                                                                                                                                                                      DB 1;
                                                                                                                                                                                  Length
                                                                                                                          Indels
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RESULT 850
ADP78301/c
ID 78301/c
ID 7878301/c
ID 7878301/
RESULT 851
ADP10746
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                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibite the expression of GFAT. Specifically claimed are antisense oligonucleotides capable of modulating the expression of GFAT, and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'-MOE wings and a deoxy gap, these oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New compounds, particularly antigense oligonucleotides targeted to a nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase (GFAT), for treating diabetes, a cardiovascular or neurologic disorder, ischemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 2100; 175pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-OCT-2003; 2003WO-US033332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GFAT; Antidiabetic; Cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chimeric phosphorothioate oligonucleotide #2100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                  1252
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                                                                                                                                                                                  TTTTGTTTTTTAATCAGA 1269
                                                                                                                          TTTTGTTTTTAAATCAAA 3
                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "
17. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= b
/mod_base= other
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                                                                                                                                                                                                                                                                                                                                                                 A; 1 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                          88.98;
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                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                          Score 14.8; DB 1; Length 20; Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                 Mismatches
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RESULT 852
ADP20152
ID ADP201
XX
AC ADP201
XX
AC ADP201
XX
DT 26-AUG
DT 26-AUG
XX
Nuclei
XX
Nuclei
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                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                   rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in a xenotransplant rejection or mechanical organ replacement rejection, in a individual. The method is also useful in assessing the immune status of an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lupus, inflammatory bowel diseases, multiple sclerosis, HIV/AIDS or viral, bacterial or fungal infection. The present sequence represents a primer for a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft rejection and other disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to diagnosing or monitoring transplant rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transplant rejection; immune system; rheumatoid arthritis; lupus; inflammatory bowel disease; multiple sclerosis; HIV; AIDS; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Set 1 left PCR primer for marker probe
                                 Nucleic acid
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20-DEC-2002; 2002US-00325899.
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                                                                 26-AUG-2004
                                                                                                 ADP20152;
                                                                                                                               ADP20152 standard;
                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 0 A; 3 C; 8 G; 9 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                             545 TGTGGGTGCGTGCGTG 562
                                                                                                                                                                                                                                                                             16;
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                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                             Conservative
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                                 detection method linking oligonucleotide #66.
                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fry K, Woodward R,
                                                                                                                                DNA;
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                                                               entry)
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                                                                                                                                                                                                                                                                                           1.1%;
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                                                                                                                                ВP
                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                           Score 14.8; DB 1
Pred. No. 4.4e+02
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                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                             2
                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                               0
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Nucleic acid

detection;

nanoparticle-oligonucleotide conjugate;

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RESULT 853
ADP20137
ID ADP201
XX
AC ADP201
XX
DT 26-AUC
XX
DT 26-AUC
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                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method of detecting a nucleic acid with at CC least two portions by providing a type of nanoparticle-oligonucleotide CC conjugate, contacting the nucleic acid and nanoparticles to allow CC hybridisation of the oligonucleotides with the two or more portions of CC the nucleic acid and observing a detectable change brought about by CC hybridisation. The oligonucleotides have a sequence complementary to the GE sequence of at least two portions of the nucleic acid. Hybridisation of CC in a detectable change. The method is used for detection and separation CC in a detectable change. The method is used for detection and separation CC of nucleic acids (e.g. viral DNA, a gene associated with a disease, bacterial DNA, fungal DNA, synthetic DNA, structurally-modified DNA, DNA CC from biological sources or PCR products) for diagnosis of various CC diseases (such as genetic diseases, bacterial infections and viral CC monitoring gene therapy. This sequencing, paternity testing and CC coligonucleotide of the invention.
                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nanoparticles useful for detection and separation of nucleic acids genes associated with disease, in a diagnostic assay, comprise seve oligonucleotides attached to them.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-JUN-1999;
13-JAN-2000;
26-APR-2000;
Nucleic acid detection; nanoparticle-oligonucleotide conjugate;
                             Nucleic acid detection method linking oligonucleotide #54.
                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-NOV-2003; 2003US-00716829.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUN-2004
                                                                26-AUG-2004
                                                                                                                           ADP20137 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-JUL-1997;
29-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US2004110220-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genetic disease; bacterial infection; viral infection; forensic; DNA sequencing; paternity testing; linking oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-JUN-2000;
26-JUN-2000;
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                                                                                                                                                                                                                                       1520 AAAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                                                                                                        16;
                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24; SEQ ID NO 70; 142pp; English.
                                                                                                                                                                                                           AAAAAAAAAAAAAAAAA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Letsinger RL,
Garimella V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-00240755.
99US-00344667.
99US-0031764099.
2000US-0200161P.
2000US-0203830.
2000US-02139069.
2001US-00760500.
                                                                                                                                                                                                                                                                      Conservative
                                                              (first entry)
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                                                                                                                                                                                                                                                                                   1.1%;
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                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                     Score 14.8; DB 1; Length 20; Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                   0 Other;
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                                                                                                                                                                                                                                                                      0;
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RESULT 854 ADP69379/c

ADP69379 standard; DNA;

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EXEXEXEX

09-SEP-2004

(first entry)

ADP69379;

human; antisense oligonucleotide; mitochondrial membrane; Human mitoNEET-specific antisense oligonucleotide #273.

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The invention relates to a method of detecting a nucleic acid with at CC least two portions by providing a type of nanoparticle-oligonucleotide cconjugate, contacting the nucleic acid and nanoparticles to allow CC hybridisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by CC hybridisation. The oligonucleotides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides have a sequence complementary to the CC in a detectable change. The method is used for detection and separation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. The method is used for detection and separation CC in nucleic acids (e.g. viral DNA, a gene associated with a disease, CC bacterial DNA, fungal DNA, synthetic DNA, structurally-modified DNA, DNA CC from biological sources or PCR products) for diagnosis of various of diseases (such as genetic diseases, bacterial infections and viral CC infections) and for forensics, DNA sequencing, paternity testing and consistoring gene thorapy. This sequence represents a linking
                                                                                                           Matches
                                                                                                                                                             Query Match
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25-JUN-1999;
13-JAN-2000;
26-APR-2000;
                                                                                                                                                                                                                    Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0
                                                                                                                                                                                                                                                                           oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Namoparticles useful for detection and separation of nucleic acids e.g. genes associated with disease, in a diagnostic assay, comprise several oligonucleotides attached to them.
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DNA sequencing; paternity testing; linking oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example
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26-JUN-2000;
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                                                                                                           16;
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                                                                                                                                     Similarity
AAAAAAAAAAAAAAAAA
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Garimella V, Li Z;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; 2000US-0176409P.
; 2000US-0200161P.
; 2000US-00603830.
; 2000US-0213906P.
; 2001US-00760500.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 55; 142pp; English.
                                                                                                           Conservative
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99US-00344667.
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                                                                                                                             1.1%;
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18
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                                                                                                                                  Score 14.8; DB 1; Length Pred. No. 4.4e+02;
                                                                                                           Mismatches
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                                                                                                                                                                                                                       0 Other;
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RESULT 855
ADF69506/c
ID ADP695
XX ADP695
XX ADP695
XX O9-SEP
XX Human;
XX human;
XX insuli
XX neurol
XX neurol
XX neurol
XX neurol
XX Homo s
XX Homo s
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XX PD 24-JUN
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention comprises antisense oligonucleotides that are targeted to the nucleic acids encoding a family of human proteins from mitochondrial membranes, which bind insulin sensitising, antidiabetic thiazolodinediones (referred to as: mitoNEET). The antisense oligonucleotides of the invention are useful for modulating mitoNEET expression and for treating diseases or conditions associated with mitoNEET, such as: diabetes, immunological disorders, cardiovascular disorders including hypertension, neurological disorders, and sechaemia/reperfusion injuries. The present DNA sequence represents a mitoNEET specific antisense oligonucleotide of the invention. NOTE: The present sequence is a 2'-methoxyethyl (2'-MOE) gapmer with a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e.g.
                                                                                                                     human; antisense oligonucleotide; mitochondrial membrane; insulin sensitising antidiabetic thiazolodinediones; mitoNEET; diabetes; immunological disorder; cardiovascular disorder; including hypertension; neurological disorders; ischaemia; reperfusion; ss; 2'-methoxyethyl gapmer; 2'-MOE gapmer; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004053060-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                      WO2004053060-A2
                                                                                                                                                                                                                                Human mitoNEET-specific
                                                                                                                                                                                                                                                                 09-SEP-2004
                                                                                                                                                                                                                                                                                                   ADP69506
                                                                                                                                                                                                                                                                                                                                      ADP69506 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 3 A; 0 C; 1 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-468836/44
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                insulin sensitising antidiabetic thiazolodinediones; mitoNEET; diabetes; immunological disorder; cardiovascular disorder; si; neurological disorders; ischaemia; reperfusion; ss; 2'-methoxyethyl gapmer; 2'-MOE gapmer; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense oligonucleotides encoding mitoNEET, useful for modulating ONEET expression or for treating diseases associated with mitoNEET, diabetes, immunological disorders or cardiovascular disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1514 TTAATTAAAAAAAAAAAA 1531
                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                           19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; SEQ ID NO 273; 226pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16;
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                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                      DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.1%;
                                                                                                                                                                                                                              antisense oligonucleotide #400.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.8;
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ADQ80728/c
ID ADQ807
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AC ADQ807
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AC ADQ807
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PORCIN
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ANOFEC
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New antisense oligonucleotides encoding mitoNEET, useful for modulating mitoNEET expression or for treating diseases associated with mitoNEET, e.g. diabetes, immunological disorders or cardiovascular disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Anorectic; Antidiabetic; Muscular; Gene Therapy; CpG island; IGF2 gene intron 3; muscle mass; fat deposition; teat number muscle deficiency; diabetes; PCR; primer; ss; pig.
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  Selecting an animal for desired genotypic or potential phenotypic
                                                                                                                                                           (UYLI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sus scrofa.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention comprises antisense oligonucleotides that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-DEC-2002;
                                                        WPI; 2004-501307/48
                                                                                                                                                                                                                                                                                                                 10-JAN-2003; 2003EP-00075091
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                                                                                                        Andersson L,
                                                                                                                                                                                                                                                              10-JAN-2003; 2003EP-00075091
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                                                                                                                                                           ) UNIV LIEGE.
) MELICA HB.
) GENTEC BV.
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                                                                                                           Andersson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.1%;
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                                                                                                        Georges
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14.8; DB 1
Pred. No. 4.4e+02
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                                                                                                           Buys
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     obesity;
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properties such as muscle mass and/or fat deposition, comprises testing for a single nucleotide polymorphism in intron 3 of the IGF2 gene.

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RESULT 857
ADP99304/c
ID ADP993
XX ADP993
XX ADP993
XX SCF/ 8
XW SCF/ 8
XW SCF/ 8
XW My3los
XW My3los
XW TREFFAC
XW TREFF
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Waplastic anaemia; paroxysmal nocturnal haemoglobinuria; myelofibrosis;

Waylosclerosis; osteopetrosis; metastatic carcinoma; acute leukaemia;

Waltiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease;

Waltemann-Pick disease; Letterer-Siwe disease;

Waltemann-Pick disease; Fulminating the syndrome;

Waltemann-Pick disease; Fulminating septicaemia; malaria;

Waltemann-Pick disease; Tulminating septicaemia;

Waltemann-Pick disease;

Waltemann-Pick 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local 9
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                                                                                            US6759215-B1
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ute leukaemia;
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AAAAAAAAAAAAAAAAAA 1 AAAAAAAAAAAGTAAAA 1537 Matches

Similarity

1.1%;

Score 14.8; DB 1 Pred. No. 4.4e+02 Mismatches

DB 1;

Length

0

Gaps

0;

0;

Sequence

20

BP;

0 ?

1 C; 1 G; 18 T; 0 U; 0 Other;

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Coperatively linked to an expression control sequence, and isolating the Coperatively linked to an expression control sequence, and isolating the Corrisponded or transfected with an expression construct comprising a CC vertebrate SCF polypeptide-encoding DNA operatively linked to a CC heterologous expression regulatory sequence, permitting the expression of CC the vertebrate SCF polypeptide in the host cell. Disclosed as new are rat CC and human mucleic acids encoding SCF, SCF protein fragments. The DNA CC sequences are useful for effecting the large scale synthesis of SCF by a CC variety of recombinant techniques or for generating new and useful viral CC and circular plasmid DNA vectors, new and useful viral CC and circular plasmid DNA vectors, new and useful transformed and circular plasmid products. The DNA sequences are also useful viral CC and circular plasmid products. The DNA sequences are also useful viral CC and tis related products. The DNA sequences are also useful viral CC protein synthesis, in genetic therapy in humans and other mammals, and CC protein synthesis, in genetic therapy in humans and other mammals, and CC paroxymmal nocturnal haemoglobinuria, myelofibrosis, multiple myeloma, CC useful for treating haematopoietic disorders, e.g., aplastic anaemia, CC paroxymmal nocturnal haemoglobinuria, myelofibrosis, multiple myeloma, CC disorders such as piebaldism and vitamin B 12 and folit acid deficiency, Diamond Blackfan anaemia, Di Guglielmo CC syndrome, congestive splenomegaly, Kala awar, sarcoidosis, primary consisting neurological damage, infertility states, intestinal damage consisting neurological damage, infertility states, intestinal damage consistence is a universal SCF pCR primer used in the isolation of SCF DNA.
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24-AUG-1990;
01-OCT-1990;
10-APR-1991;
25-NOV-1992;
21-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Preparing a human stem cell factor (SCF) polypeptide, useful for treating hematopoietic disorders, e.g., aplastic anemia, comprises growing host cells transformed or transfected with DNA encoding a human SCF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to preparing a (vertebrate) human stem cell fa
(SCF) polypeptide comprising growing host cells transformed or
transfected with DNA encoding a human SCF that stimulates growth of
haematopoietic progenitor cells under nutrient conditions, the DNA h
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example
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92US-00982255.
93US-00172329.
95US-00449182.
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90US-00573616.
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RESULT 858
ADP99302/c
ID ADP993
The invention relates to preparing a (vertebrate) human stem cell factor (C (SCF) polypeptide comprising growing host cells transformed or transfected with DNA encoding a human SCF that stimulates growth of the comprisite progenitor cells under nutrient conditions, the DNA being CC operatively linked to an expression control sequence, and isolating the CC operatively linked to an expression control sequence, and isolating the CC polypeptide produced. Also included is a recombinant host cell the control sequence, permitting the control sequence programment of the control sequence, permitting the expression of the vertebrate SCF polypeptide encoding DNA operatively linked to a control sequence, permitting the expression of the vertebrate SCF polypeptide in the host cell. Disclosed as new are rated and human nucleic acids encoding SCF, SCF proteins from a number of other comminate and recombinantly expressed SCF proteins from a number of other comminate and recombinantly expressed SCF protein fragments. The DNA sequences are useful for effecting the large scale synthesis of SCF by a cariety of recombinant techniques or for generating new and useful viral and circular plasmid DNA vectors, new and useful transformed and croamer of the polypeptide products. The DNA sequences are also useful as control of the protession of such host cells and new and useful as control of the protession of the polypeptide products. The DNA sequences are also useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-APR-1991;
25-NOV-1992;
21-DEC-1993;
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11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   refractory erythroblastic anaemia; Di Guglielmo syndrome; congestive splenomegaly; Kala awar; sarcoidosis; primary splenic pancytopaenia; miliary tuberculosis; disseminated fungus disease; Fulminating septicaemia; malaria; vitamin Bl2 deficiency; folic acid deficiency; pyridoxine deficiency; Diamond Blackfan anaemia; hypopigmentation disorder; piebaldism; vitiligo; neurological damage; infertility; intestinal damage; irradiation; chemotherapy; AIDS; haematopoletic recovery; acute blood loss; neoplasm; cancer; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease; Niemann-Pick disease; Letterer-Siwe disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               aplastic anaemia; paroxysmal nocturnal haemoglobinuria; myelofibrosis; myelosclerosis; osteopetrosis; metastatic carcinoma; acute leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Preparing a human stem cell factor (SCF) polypeptide, useful for treating hematopoietic disorders, e.g., aplastic anemia, comprises growing host cells transformed or transfected with DNA encoding a human SCF.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SCF; stem cell factor; gene therapy; haematopoietic progenitor cell;
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90US-00573616.
90US-00589701.
91US-00684535.
92US-00982255.
93US-00172329.
95US-00449182.
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human genomic
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SCF,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 16
                             A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                    WPI; 1995-018287/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                                                                                   Disclosure; Page 7; 11pp; Japanese.
                                                                                                                                                                                                                    by digestion
                                                                                                                                                                                                                                   Analysis of cDNA and
                                                                                                                                                                                                                                                                                                                                                                       16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ75707;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ75707
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                                                                                                                                                                                                                                                                                                                                      16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene expression; reverse transcription; primer; cDNA;
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                                                                                                                                                                                                                                   amplification of mRNA followed
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RESULT 860
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XX PCR; 8
Micros
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                                                                                                                                                                                                                                                                                                    The sequence is that of a bovine microsatellite sequence obtd. by CC screening a library of bovine MboI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 CC clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (T6) >9 microsatellites of the bovine genome is estimated at >100, 000. The sequence information CC in the bovine genome is estimated at >100, 000. The sequence unformation CC for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequences upstream and CC downstream of the microsatellite sequence were used to generate the required PCR primers for in vitro amplification of the corresp.

CC microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of correct pince in the program of the program of traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Table 7; Page 245; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polymorphic bovine DNA markers - used in genetic identification, mapping, and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-AUG-1992.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Microsatellite sequence from clone TGLA2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
02-FEB-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Georges M, Massey JM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GENM-) GENMARK.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1519 TAAAAAAAAAAAGTAAA 1536
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                                                                                                                                                                                                                           BP; 0 A;
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Conservative
                                                                                                              Conservative
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(first entry)
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                                                                                                                                                                                                                           1 C; 10
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                                                                                                            Score 14.8; D
Pred. No. 4.2e
0; Mismatches
                                                                                                                                                                                                                        G; 10 T; 0 U; 0 Other;
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Pred. No. 4.2e+02;
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                                                                                                                                         4.2e+02
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                                                                                                                                                                      DB 1;
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                                                                                                                                                                Length 21;
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RESULT

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AAQ75702/c
ID AAQ75702 standard; DNA; 21 BP.
XX AAQ75702;
XX AAQ75702;
XX AAQ75702;
XX AAQ75702;
XX AAQ75702;
XX Reverse transcription primer use XX Analysis; gene expression; rever XX aggregate; restriction enzyme; s XX Synthetic.
XX JP06303997-A.
XX JP0630303997-A.
XX JP06303997-A.
XX JP06303
RESULT 862
AAQ75671/c
ID AAQ756
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Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Analysis; gene expression; reverse aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analysis of cDNA and gene expression – by amplification of mRNA followed by digestion with restriction enzymes.
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         16-APR-1993;
                                                                                                 16-APR-1993;
                                                                                                                                                                                             01-NOV-1994.
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                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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         93JP-00112515
                                                                                                      93JP-00112515.
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Pred. No. 4.2e+02;
0; Mismatches 2;
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XX JP0630
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XX AAQ76
PD 01-NOV
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Best Local Similarity
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                16-APR-1993;
                                          16-APR-1993;
                                                                      01-NOV-1994
                                                                                                                          Synthetic
                                                                                                                                                   aggregate;
                                                                                                                                                               Analysis; gene expression; reverse transcription;
                                                                                                                                                                                         Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                     04-AUG-1995
                                                                                                                                                                                                                                               AAQ75688;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 2 A; 1 C; 0 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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aggregate; restriction enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Reverse transcription primer used
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                                                                                                                                                                                                                                                                          standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                be used
                                                                                                                                                    restriction enzyme; ss
                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                           93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             to analyse gene expression rapidly
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Pred. No. 4.2e+02;
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RESULT 870
AAQ75715/c
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XX JP0630
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Best Local :
                                                                   A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The
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                                                                                                                                                                                                                                                                                                          Analysis of cDNA and gene expression -
by digestion with restriction enzymes.
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  Sequence
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                                                  method can
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-APR-1993;
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    21 BP; 1 A; 1 C; 2 G; 17 T; 0 U; 0 Other;
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                                                be used to analyse gene expression rapidly
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Pred. No. 4.2e+02;
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Query Match

Score 14.8;

DB

1:

Length 21;

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RESULT 871
AAQ75686/c
ID AAQ756
XX AAQ76

AAQ75689/C
ID AAQ756
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Matches 16; Conserv
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                     Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                           Reverse transcription primer used in cDNA analysis technique.
                                                                                                                 04-AUG-1995
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                                                                                                                                                                                              AAQ75689 standard;
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aggregate; restriction enzyme;
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Pred. No. 4.2e+02;
0; Mismatches 2;
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RESULT 873
AAQ75703/c
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Best Local
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                                      Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reverse transcription primer used in cDNA analysis technique.
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Disclosure; Page 7; 11pp; Japanese
                                                                                                     WPI; 1995-018287/03.
                                                                                                                                                                                       16-APR-1993;
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A method for the analysis of cDNA comprises double-stranded cDNAs by using an aggregate

(a) preparing an aggregate of mRNAs and a plural type

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RESULT 874
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ID AAQ75672 standard; DNA;
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 7; 11pp; Japanese.
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                                                                                                                                                                                                                                                              ch 1.1%; 1 Similarity 88.9%; 16; Conservative
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0; Mismatches
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Pred. No. 4.2e+02;
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                                                                                                                                                                                                                                                                                                                       DB 1;
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RESULT 876
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Best Local (
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                                                                                                                                                                                                                                                         AAQ75706 standard;
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                        16-APR-1993;
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                                                                                                                                             aggregate;
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(NITE ) NIPPON
                                               16-APR-1993;
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                                                                                                                                                                                                                                                                                                                        18 TAAAAAAAAAAAAAAA 1
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                                                                                                                                                          gene expression; reverse
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                                                                                                                                                restriction enzyme;
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TELEGRAPH & TELEPHONE CORP
                        93JP-00112515
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                                                93JP-00112515
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Pred.
                                                                                                                                                88.
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No. 4.2e+02;
                                                                                                                                                           transcription; primer; cDNA;
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AAQ75685/c
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XX
AC AAQ756
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AC AAQ756
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Revers
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                                                                                                                                                                                                                                                                                            gene expression; reverse transcription;
; restriction enzyme; ss.
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ilarity 88.9%;
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Pred. No. 4.2e+02;
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RESULT 88
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                                                                                                              Sequence
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by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                       Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                      16-APR-1993;
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                          1519 ТААЛАЛАЛАЛАЛАСТАЛА 1536
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Pred. No. 4.2e+02;
                                                       Score 14.8; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 2
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Pred. No. 4.2e+02;
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Query Match Best Local S Matches 16

l Similarity 88. 16; Conservative

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Score 14.8; DB 1; Pred. No. 4.2e+02;

DB 1;

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Mismatches

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Best Local &
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                                                                             aggregate;
                                                                                   Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                 Reverse transcription
                                                                                                                                                                                                                   04-AUG-1995
                                                                                                                                                                                                                                                                                                                             AAQ75701 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Reverse transcription primer used
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Pred. No. 4.2e
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A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse
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                                                                                                       Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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Pred. No. 4.2e+02;
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Matches 16
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                                                                                             The sequences represented by, AAQ90390-90401 are synthetic DNA probes containing 3' ribonucleoside termini. The sequences shown in AAQ90402-15 are synthetic DNA probes with 5' amino termini. These sequences were specific for the polymorphisms of HLA gene dQa. The sequences were used in the device of the invention. This is a self-addressable electronic device (SAED) that can be used to carry out multi-step and multiplex reactions, such as nucleic acid hybridisations. The advantages of this method are that these reactions can be carried out with complete and precise electronic control, and that the rate, specificity and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                        Sequence
                                                                                                                                                                                                                                        Example 1; Page 40; 86pp; English.
                                                                                                                                                                                                                                                                                   New self-addressable electronic multiplex reactions such as DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CP-1 (synthetic DNA probe with 3'ribonucleoside terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ90391;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ90391 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                  sensitivity
                                                                                                                                                                                                                                                                                                                                                                                    (NANO-) NANOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                    bio:polymer synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1519 ТАЛАЛАЛАЛАЛАГТАЛА 1536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          \mathsf{HLA}_i dQa; 3' ribonucleoside; self-addressable electronic device; hybridisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18
  l Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
16; Conserv
                                                       21 BP; 20 A; 0 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TANANANAAAAAAAA 1
                                                                                  of these reactions are greatly improved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry
                                                                                                                                                                                                                                                                                                                                                                                                              93US-00146504
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag=  a
/note= "3′ ribonucleoside terminal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.1%;
              1.1%;
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Score 14.8; DB 1; Length 21; Pred. No. 4.2e+02; 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No.
                                                                                                                                                                                                                                                                                   devices - used for multi-step and
hybridisation(s), clinical diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                    at micro-locations
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  Gaps
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RESULT 890 AAV35395 ID AAV353 XX

AAV35395 standard; DNA; 21

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                                                   Matches
                                                                           Query Match
                                                                                                                         are used in the construction of the electronically self- addressable device (ED) of the invention. The ED comprises a substrate, an electrode or opt. a number of electrodes supported by the substrate, a current source operatively connected to the electrode and an attachment layer adjacent to the electrode which is permeable to a counterion but not permeable to a molecule capable of insulating or binding a macromolecule. The attachment layer is capable of attaching a macromolecule. The ED is used for genetic typing and comprises a number of electronically addressable locations each comprising an electrode, and a binding entity, such as one of these probes, attached to each of the locations capable of detecting the presence of a genetic sequence
                                                                                                                                                                                                                                                                                                                                Electronically self-addressable device - used for electronic control of, e.g. nucleic acid hybridisation.
                                                                                                      Sequence 21 BP;
                                                                                                                                                                                                                                                                                                       Example 1; Page 60; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                       WPI; 1996-097582/10.
                                                                                                                                                                                                                                                                                                                                                                                                 Heller MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   attachment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Electronically self-addressable device; ED; electrode; current source;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT10743 standard; RNA; 21 BP
                                                                                                                                                                                                                                                                             The sequences given in AAT10742-67 are synthetic oligonucleotides which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JAN-1996
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                                                                Local
                1520 AAAAAAAAAAAAGTAAAA 1537
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                                                   16;
                                                                 Similarity
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 AAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 layer; permeable; counterion; genetic typing; probe;
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Έ
                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                      20 A; 0 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        probe, CP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                   Evans
                                                             1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a
"3'-ribonucleoside terminus"
                                                                                                                                                                                                                                                                                                                                                                                                 GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18
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                                                   0;
                                                 Pred. No. 4.2
                                                                Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                 Sosnowski
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                                                                             DB 1;
                                                                           Length 21;
                                                     Indels
                                                   <u>,</u>
                                                   Gaps
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                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                             Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                            Human polymorphic region 592.
                                                                                                   30-NOV-1999
                                                                                                                         AAZ26403;
                                                                                                                                              AAZ26403
                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                            protecting against infection with
                                                                                                                                                                                                                                                                                                                                                                                                                        such as HIV-1 or HIV-2, fused to a DNA fragment containing an epitope region (30-300 bases in length) of a retroviral gene other than the g
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV35388-V35414 are primers used in a method which results in a retrovirus recombinant vaccinia virus comprising of a gag gene from a retrovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 66; 84pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Recombinant vaccinia virus containing fusion HIB gag gene production in host cells of gag protein for use as vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kojima A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-NOV-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         vaccines; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypervariable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV-1 gag protein DNA primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-OCT-1998
                                                                                                                                                                                                                        1520
                                                                                                                                                                                                                                              16;
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                                                                                                                                                                                                                                                        Similarity
                                                                                                                                              standard; DNA; 21
                                                                                                                                                                                                   AAAAAAAAAAAAAAAAAA 20
                                                                                                                                                                                                                      дадададададатадад 1537
                                                                                                                                                                                                                                                                                       21 BP; 19 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kurata T,
                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   region; ENV protein; vaccinia virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96JP-00323412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97WO-JP004216
                                                                                                                                                                                                                                                       1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protection; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yasuda
                                                                                                                                              BP
                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                       Score 14.8;
Pred. No. 4.
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                                                                                                                                                                                                                                                      4.2e+02;
                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene; retrovirus;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 16
                                                             Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; athlele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                    Human polymorphic region 415.
                                                                                                                                                                                                                       30-NOV-1999
                                                                                                                                                                                                                                                         AAZ26226;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 12 A; 2 C; 0 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 7; 605pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                                                                                                                                          AAZ26226 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Housman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                   1514 TTAATTAAAAAAAAAAA 1531
                                                                                                                                                                                                                                                                                                                                                                                                                                               16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphic sites described in the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                             TTTTTAAAAAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.1%;
                                                                                                                                                                                                                                                                                          21
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                                                                                                                                                                                                                                                                                                                                                                             21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21;
                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Homo sapiens

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RESULT 893
AAX81302
ID AAX813
XX AAX813
XX AAX813
XX CAAX813
XX IO-POT 20-AUG
XX IO-POE
XX Microe
XX Microe
XX IO-POE
XX Synthe
XX Synthe
XX Synthe
XX W100-POE
XX W09929
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC This invention describes a novel method for identifying an inhibitor CC potentially useful for treatment of cancer, where the inhibitor is active CC on a gene vital for cell growth or viability, and where the gene is CC subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is CC used for preventing the development of cancer in a patient having a CC used for preventing the development of cancer in a patient having a CC precancerous condition, by administering to the patient a first allele CC specific inhibitor (ASI) targeted to an allele of a first sesential gene present in cells of the precancerous condition, where the normal somatic CC cells of the patient are heterozygous for the first gene, the inhibitor CC cancer in a population and tragets only one allelic forms of the gene CC used in the diagnosis, prevention and treatment of LOH disorders, e.g. CC cancers, atheroscleroic plaques, premalignant metaplastic or dysplastic CC lesions, benign tumours, endometriosis, polycystic kidney disease, and CC graft versus host disease. The method can also be used to remove CC malignant cells from bone marrow transplants. AAZZS812-Z26825 represent CC human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local 9
                                                                                                                                                                                                   Microelectronic device; multi-step reaction; microscopic format; ion-permeable permeation layer; electrode; electrical control; transport; attachment; binding; DNA/RNA hybrid; probe; ss.
                                                                                                                                                                                                                                                                                3' ribonucleoside oligonucleotide probe CP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                         20-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                 AAX81302 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 2 A; 6 C; 12 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Fig 7, 605pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-MAR-1997;
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                                                        W09929711-A1
                                                                                                               misc_RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GAGGGGCGGCCGCCGCC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; DB 1;
Pred. No. 4.2e+02;
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17-JUN-1999

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RESULT 894
AAV71751/c
ID AAV717
XX AAV717
XX Human
DT 15-MAR
XX HIV re
KW HIV re
KW P30 pr
XX Synthe
OS Homo s
XX HOMO 8
DF 17-SEP
XX 12-MAR
XX 11S-MAR
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XX (INSP
PA (INSP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            microelectronic device which is designed to actively carry out and control multi-step and multiplex molecular biological reactions in microscopic formats. A key aspect of this inventions is played by the ion permeable permeation layer which overlies the electrode. This permeation control multi-step and multiplex molecular biological reactions. This permeation layer allows attachment of nucleic acids to permit immobilization but also separates the attached oligonucleotides and hybridizet target DNA sequences from the highly reactive electrochemical environment generated control tentions and fabricated to actively carry out and control reactions such as nucleic acid hybridizations, antibody/antigen reactions such as nucleic acid hybridizations, antibody/antigen reactions as such as nucleic acids and polypeptides, to specific microlocations. The device can subsequently control the transport and reaction of analytes or reactants at the addressed specific microlocations. The device is able to concentrate analytes and reactants, remove non-specifically bound molecules, provide stringency control for DNA hybridization reactions and improve the detection of analytes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                             HIV receptor; V3 loop; human immunodeficiency virus; P30 protein; PHAPI; infection; therapy; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and
                                                  12-MAR-1997;
                                                                                 12-MAR-1998;
                                                                                                                 17-SEP-1998
                                                                                                                                                 WO9840480-A1
                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                              Human V3 loop
                                                                                                                                                                                                                                                                                                               15-MAR-1999
                                                                                                                                                                                                                                                                                                                                              AAV71751;
                                                                                                                                                                                                                                                                                                                                                                              AAV71751 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The specification describes a self-addressable, self-assembling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 89; 179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-385567/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sosnowski RG, Butler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-DEC-1998;
   (INSP ) INST PASTEUR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NANO-) NANOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    microelectronic device designed to carry multiplex molecular biological reactions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1520 AAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                             sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence represents a probe used to exemplify the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAAAAAA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                              HIV receptor P30/PHAPI sense PCR primer.
                                                  97US-0040969P
                                                                                  98WO-EP001409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-00986065.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US025475.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 A; 0 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nerenberg MI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     t and control multi-step microscopic format.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Heller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                retrovirus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Edman CF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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This oligonucletide is complementary to a portion of DNA sequence (see AAV71743) coding for the p30/PHAPI (see AAW84053) of the newly identified V3 loop HIV receptor. It is used as a sense primer, together with an cantisense primer (see AAV71752) in a PCR amplification of p30/PHAPI reverse-transcribed mRNA. The V3 loop HIV receptor consists of an association of 3 proteins, named P95/mucleolin, P40/PHAPI and P30/PHAPI (see AAW84052-54). A method for screening molecules that modulate the expression of the receptor comprises: cultivating cells transfected with a nucleotide sequence encoding P95/mucleolin, P30/PHAPI or P30/PHAPII, C2 placed under the control of its own promoter; bringing the cells into contact with a test molecule; and quantifying expression of the P95/mucleolin, P30/PHAPI or P30/PHAPI o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                      WPI; 1999-479058/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated V3 loop HIV receptor - comprises P95/nucleolin, P40/PHAPII and P30/PHAPI proteins, used to develop products for the treatment and prevention of HIV infection.
                                                                                                                             Brooks-Wilson AR,
Miller A, North
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 1 A; 10 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hovanessian Briand J, (
                                                                                                                                                                                                                                21-JAN-1998;
                                                                                                                                                                                                                                                                           21-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                       WO9937809-A1
                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                       therapeutic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ASTH1; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic fragment in region 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ18416 standard; DNA; 21 BP
                                                                                                                                                                                        (AXYS-) AXYS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   427 GCGGCTGCGGCGCGGCG 444
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Guichard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 48; 267pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                     a; human; chromosome 11p;
immunogen; polymorphism;
                                                                                                                                                                                                                                98WO-US001260
                                                                                                                                                                                                                                                                        98WO-US001260.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Callebaut hard G;
                                                                                                                               3
                                                                                                                                                  Buckler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Krust B,
                                                                                                                                                  Cardon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to ASTHLJ.
                                                                                                                                                                                                                                                                                                                                                                                                                                       ASTH1I; ASTHIJ; genetic locus; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jacotot E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .2e+02
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                                                                                                                                                  Carey
                                                                                                                                                  ₽H,
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                                                                                                                                                    Galvin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Mammalian asthma related genes,

useful for diagnosis

of a predisposition

of asthma

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RESULT 896
AAX26973/c
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Best Local S
Matches 16
       The present primer was used to reverse transcribe RNA encoding a human mammary-specific protein, designated mammaglobin. The specification describes a protein comprising a mammaglobin antigen that is recognized by B and/or Tc cells specific for the natural, secreted and glycosylated form of mammaglobin polypeptide. This protein, or recombinant vectors that express it, are used in vaccines for treating mammaglobin-expressity cancers, specifically of the breast. Such cancers can also be treated using autologous tumor lymphocytes activated ex vivo with an mammaglobin is elevated in 77% of stage I primary breast cancers, so it represents a marker useful for diagnosis of this disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mapped to human chromosome Ip. ASTHII and ASTHIJ are genes present within the locus, located close to each other on human chromosome 11p, and have similar patterns of expression, and common sequence motifs. The ASTHI genes and fragments, encoded protein, genomic regulatory regions and anti-ASTHI antibodies are useful in the identification of individuals predisposed to development of asthma, and for the modulation of gene activity in vivo for prophylactic and therapeutic purposes. The ASTHI protein is useful as an immunogen to raise specific antibodies, in drug screening for compositions that mimic or modulate ASTHI activity or expression, including altered forms of ASTHI protein, and as a therapeutic. Sequences AAZI8366-Z18509 represent polymorphisms in the ASTHII and ASTHIJ genes
                                                                                                                                                                                                                              Example 2; Page 55; 60pp; English.
                                                                                                                                                                                                                                                                 Mammaglobin, secreted protein overexpressed
                                                                                                                                                                                                                                                                                                                                       Watson MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; mammary-specific protein; mammaglobin; antigen; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX26973;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX26973 standard; cDNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 0 A; 9 C; 1 G; 10 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention identifies a genetic locus ASTH1, associated with asthma, mapped to human chromosome 11p. ASTH1I and ASTHIJ are genes present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 62; 195pp; English
                                                                                                                                                                                                                                                                                                    WPI; 1999-244021/20.
                                                                                                                                                                                                                                                                                                                                                                                                            18-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                             18-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9914230-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        aucorogous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mammaglobin-expressing cancer; breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                          (UNIW ) UNIV WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1225 CTCTAGCTCTTAGCTTCCTC 1244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tumor lymphocyte;
                                                                                                                                                                                                                                                                                                                                     Fleming TP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              to reverse transcribe mammaglobin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                             97US-00933149.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diagnosis; marker; primer;
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Pred. No. 4.2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                 in breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA
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                                                                                                                                                                                                                                                                    cancer
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ID AAZA43
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XX Protei
XX AATIAN
KW Prophy
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RESULT 89
AAA80318
ID AAA6
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                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 16
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes novel purified aptameric oligonucleotides which have antimicrobial, cytostatic and immunosuppressive activity. The oligonucleotides are useful for binding to and preventing or inhibiting the biological function of a protein kinase or a target molecule and for detecting the presence or absence of a target molecule in biological samples. The oligonucleotides are also useful for prophylactic and therapeutic treatment of diseases such as cancer, autoimmune diseases and diseases caused by pathogenic microorganisms. This sequence represents a primer used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonuclectides inhibiting protein kinase, useful for treating diseases such as cancer and autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antimicrobial; cytostatic; immunosuppressive; protein kinase; prophylactic; therapy; treatment; cancer; autoimmune disease; pathogenic microorganism; primer; ss.
                            AAA80318
                                                                                 AAA80318
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 8; Col 27-28; 26pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-DEC-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Protein kinase inhibiting primer #12.
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                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                 standard;
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                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 0
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                                                                                 DNA;
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Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8;
                                                                                                                                                                                                                                                                                                                                                          Pred.
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                                                                                                                                                                                                                                                                                                                                                          No.
                                                                                                                                                                                                                                                                                                                                                          4.2e+02
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CC Chromosome (11p). This locus comprises the ASTHII and ASTHII genes, which CC chromosome (11p). This locus comprises the ASTHII and ASTHII genes, which CC are associated with a genetic prediposition to asthma and bronchial CC hyperreactivity. The ASTHII and ASTHII genes are oriented in opposite CC directions with the ASTHII locus, and have similar patterns of expression CC and common sequence motife. They are both expressed in trachea, lung and CC several other tissues. ASTHII and ASTHII are novel members of the ets CC family of transcription factors, which have been implicated in the ets Granily of transcription factors, which have been implicated in the CC genes known to be important in the actiology of asthma. Both ASTHII and CRASTHII mand craim mand craim a second management of asthma, as probes for identifying ASTHII related genes, CRASTHII genesion of the gene in a biological specimen, and for generating genetically modified non-human animals or site specific gene campositions that mimic or modulate activity or expression of ASTHII candiforms of these proteins; and as a CRASTHII genes or fragments thereof, encoded proteins, and as a CRASTHII genes or fragments thereof, encoded proteins, and second and therapeutic purposes. The intact ASTHII and anti-ASTHII crasting or prophylactic and thearapeutic purposes. The intact ASTHII or ASTHII crasting crasting proteins or active fragments thereof may be used to modulate or reduce bronchial hyperreactivity. Sequences AAA80260-A80261 and AAA80264-A80416 crapressent polymorphic sites within the ASTHII or ASTHII genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acids other than naturally occurring chromosomes encoding ASTH1 protein, for e.g. screening compositions that modulate expression or function of ASTH1 proteins or as diagnostics for genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   bronchial hyperreactivity; ets family; transcription splice variant; genetic predisposition; polymorphism; drug screening; prophylaxis; therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Galvin M, Mil
Brooks-Wilson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-JAN-1997;
01-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Col 41-42; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   predisposition to asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ASTH1 locus; ASTH11; ASTH1J; human; chromosome 11p; asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AXYS-) AXYS PHARM INC.
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21
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B₽;
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97US-0051432P.
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A;
9 C; 1
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G; 10 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cardon L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       site,
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 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Buckler
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Query Match Best Local : Matches 1:

Similarity

80.0%;

1.

1.1%;

Score 14.8; Pred. No. 4.

4.2e+02

DB 1;

Length 21; Indels

0

Gaps

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RESULT 899
AAF9977/C
ID AAF997
XX AAF997
AC AAF997
AC AAF997
XX DT 12-JUN
XX Vaccin
KW Vaccin
KW inmuno
KW fungal
KW infect
XX Synthe
XX 25-SEP
PR 27-SEP
RX 27-SEP
RX
RESULT 900
AAH42480/c
ID AAH42480 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acid can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           haemophilus, campylobacter, clostridium, Escherichia coli and/or haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 101; Page 56; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vaccinating against tumors, infectious diseases, allergies and using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-SEP-1999;
27-SEP-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunostimulatory nucleic acid #823.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-AUG-2000; 2000US-0227436P
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                                                                                                                                                                                                                                                                                       1 Similarity
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                                                                                                                                                                                                                        ААААААААААБТААА 1537
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                                                                                                                                                                AAAAAAAAAAAAAAAA 4
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                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                          Conservative
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99US-0156135P.
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                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                        Score 14.8;
Pred. No. 4.
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                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                           4.2e+02;
                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                  Length 21;
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                                                                                                                                                                                                                                                                                       0,
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RESULT 901
ABS78428/c
ID ABS784
XX
AC ABS784
XX
AC ABS784
XX
DT 13-DEC
XX
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                                                                                                                                                                                                                                                                                                           Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                       moletles optionally extended by a polymerase. The branched chain compounds of the invention are used in nucleic acid synthesis reaction, primer extension reaction, reverse transcription reaction of RNA into DNA, nucleic acid hybridization experiment (for identifying sequence of a nucleic acid), and nucleic acid amplification experiment (for analysing the expression pattern of genes). The compounds are also used in solid-phase enzymatic reactions. The present sequence was used in the course of the invention to produce branched chain compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Branched chain compound; nucleic acid synthesis; primer extension; reverse transcription; nucleic acid hybridization; nucleic acid amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Branched compounds useful in e.g. nucleic acid synthesis reaction comprises nucleic acid moieties optionally extended by a polymera
                       13-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schmidt W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide used to produce branched chain compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-OCT-2001
                                                               ABS78428;
                                                                                                     ABS78428 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 10; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (VBCG-) VBC GENOMICS GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The specification describes branched compounds containing nucleic acid
                                                                                                                                                                                                                                                1520 AAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                          21
                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                          Conservative
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                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note=
                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                           1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        b
"NH2-C6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "branch present"
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"NH2-C6 attached"
                                                                                                       21
                                                                                                       BP
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                                                                                                                                                                                                                                                                                                           Score 14.8;
Pred. No. 4.
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                                                                                                                                                                                                                                                                                          Mismatches
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AC ABL394
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Immunc
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis aspociated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                       ABL39404;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telanglectasia; haemophiliac joint; plaque neovascularisation; telanglectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
                                              Antibody-induced cell lysis; cancer; immunostimulatory; CD20; anglogenesis; metastasis; cytostatic; phosphorothicate backbone; ss
                                                                                                                                       16-APR-2002
                                                                                                                                                                                                         ABL39404 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 35; 276pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inhibiting angiogenesis in a subject, involves administering at antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-DEC-2000; 2000US-025534P.
                                                                                                    Immunostimulatory nucleic acid SEQ ID NO: 840.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-DEC-2001; 2001WO-US048458.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                scleroderma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumour metastasis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (COLE-) COLEY
                                                                                                                                                                                                                                                                                                                               1520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of the invention
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                                                                                                                                                                                                                                                                                                                            AAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    inhibitor; 88; angiogenesis; solid tumour growth;
                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                            1.1%;
                                                                                                                                                                                                         21
                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                  G; 21 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                                Score 14.8;
Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
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RESULT 903
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Best Local
                            Human; ds; cytochrome P450 A1; CYP4501A1; UGT2B4; MDR1; cytochrome P450 A2; CYP4501A2; cytochrome P450 O2E; CYP45002E1; LTF; adrenergic receptor beta1; ADBR1, aryl hydrocarbon; AHR; MRP3; NR112; aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS cyclooxgenase 2; COX2; diazepam binding inhibitor; DBI; haematological; epoxide hydroxylase 2; EPHX2; S-lipoxygenase activating protein; FLAP; glutathione-S-transferase; 12; GST12; histamine-N-methyl transferase; NNMT; kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, cervical cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, non-Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovarian cancer, stomach cancer, testicular cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, rhabdomyosarcoma, skin the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present sequence is an immunostimulatory oligonucleotide described exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI;
NADPH quinone oxidoreductase 2; NQO2; sulfotransferase thermolabile; STM; UDP-glucuronosyl transferase 2B7;
                                                                                                                                                                                          Human NADPH
                                                                                                                                                                                                                            23-DEC-2002
                                                                                                                                                                                                                                                                                              ABS97830 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Weiner G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                              ABS97830
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     developing cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (IOWA ) UNIV IOWA RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-JUN-2000; 2000US-0213346P
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                                                                                                                                                                                                                                                                                                                                                                                                                 1520 АЛАЛАЛАЛАЛАСТАЛАЛ 1537
                                                                                                                                                                                                                                                                                                                                                                                  21
                                                                                                                                                                                                                                                                                                                                                                                                                                                    16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                AAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                      quinone oxidoreductase 2 (NQO2) polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 0 A; 0 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 309; 312pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                        (first entry)
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1%;
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                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length
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                                                                                                                                                                                          sequence #38.
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or at risk of
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CC This invention relates to the sequence of an isolated nucleic acid compositing at least one base variation from that of a known chuman cytochrome P450 Al (CYP4501Al), cathepain S (CTSS), cytochrome P450 Al (CYP4501Al), cytochrome P450 
UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA; multidrug resistance 1; lactotransferrin; orphan nuclear receptor; multidrug resistance associated protein 3; cancer; prostate; acetylcholine muscarinic receptor; CHWR1; CHWR2; CHWR3; CHWR4; CHWR5; altered drug metabolism; cardiovascular function; colorectal tumour; central nervous system; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 16; Page 130; 714pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disorder-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-698522/75.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleotide polymorphism.
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Best Local Similarity

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Score 14.8; DB 1; Length 21; Pred. No. 4.2e+02;

Query Match

Sequence

21

B₽; DNA

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A; 9

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G; 1 T; 0 U; 0 Other;

nervous system

function. The present sequence represents

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sequence of

invention

S 밁 Query Match Best Local S Matches 16 remote diagnostic server; processing digital sample signal at remotely located database to correlate digital signal with digital information and returning report of particular condition of animal. The method is useful for assessing a condition of a performance animal preferably human, dog or camel. The condition of a performance animal preferably human, dog or camel. The condition can be an athletic ability and a condition that enhances, hinders, impedes or does not change an expected ability of the performance animal; and also normal, pre-clinical, overt progress and/or stage of disease, undiagnosed of unclassified conditions, presence of drugs, response to exercise, response to vaccines, therapies, nutritional states and response to environmental conditions. Diseases assessed by the invention include laminitis, lameless, viral or bacterial disease, epistaxis, musculoskeletal damage or disorders and joint diseases. The present sequence is a primer used to illustrate the method of the invention The invention relates to a method for assessing a condition of a performance animal. The method involves determining in sample abundance of expressed target nucleic acid; transmitting digital sample signal to Assessing condition e.g. athletic ability, stage of disease, presence of drugs, response to exercise, response to vaccines, therapies, nutritional 04-MAY-2001; 29-JUN-2001; Laminitis; viral disease; vaccine; bacterial disease; primer; epistaxis; gastritis; gastric ulcer; respiratory ailment; fracture; joint disease; Sequence 21 Disclosure; Page 46; 87pp; English. states, of WPI; 2003-120558/11. Brandon 03-MAY-2002; 2002WO-AU000553 14-NOV-2002 Unidentified. musculoskeletal damage; ss. Regular oligo dT 16-APR-2003 AAD51323; AAD51323 standard; (GENO-) GENOMICS 1520 AAAAAAAAAAAAGTAAAA 1537 Similarity performance animal involves analyzing nucleic acid BP; Conservative 2001AU-00004809. 2001US-00896941. (first entry) 0 primer used to illustrate the method RES PARTNERS Þ DNA; 1.1%; 0 C; 0 G; 21 ВÞ <u>,</u> Pred. No. 4.2 0; Mismatches Score 14.8; YTG 21 T; 0 LTD. 4.2e+02 U; 0 Other; DB 1; Length 21; 얁 the invention 0 expression Gaps

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ADB37209/C
ID ADB372
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AC ADB372
XY
AC ADB372
XX
DT 04-DEC
XX
DE Immunc
XX
C
KW ds; al
KW hypo-x
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OS Synthe
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ACH03246/c
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                      The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Croh's disease.
                                                                                                                                                                  1209/c
ADB37209 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 33; 229pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating non-allergic inflammatory diseases, such as psoriasis, allergic contact dermatitis, latex dermatitis or inflammatory b
                                        ds; allergy; asthma; poly-G nucleic acid; aerosol formulation;
hypo-responsive subject; immunostimulatory.
                                                                                                                                         ADB37209;
                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antilicer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contect dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
                                                                                                               04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAR-2001; 2001US-0279642P.
                                                                                 Immunostimulatory nucleic acid #823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-MAR-2002; 2002US-00112653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US2003050268-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunostimulatory nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACH03246;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACH03246 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BERG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (KRIE/)
                                                                                                                                                                                                                                                                   1520
                                                                                                                                                                                                                                                                                                                                                                            sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                             1 Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BERG D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KRIEG A M.
                                                                                                                                                                                                                                                       AAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                       AAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Berg DJ
                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                          1.1%;
                                                                                                                                                                   21
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                                                                                                                                                                   ВP
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                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                           Score 14.8; Di
Pred. No. 4.2e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #881.
                                                                                                                                                                                                                                                                                                          4.2e+02
                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                             0,
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ADK01309/c
XX
AC ADK01
XX
AC ADK01
XX
XX Rat |
DE Rat |
XX
KW 88;
KW 5100
XX
COS Ratt
XX
DE Rat |
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DE Rat |
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 907
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BRAT/)
(PETE/)
(FOUR/)
                                                                         Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                               DE10208794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rattus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rat DNA microarray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 17; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI;
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                                                  and constant regions
                                                                                                                                                     WPI; 2003-714082/68.
                                                                                                                                                                                                 Boekenkamp D,
                                                                                                                                                                                                                                                 (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                 28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                               28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bratzler RL,
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FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                 Dieck
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 0 C; 0 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    capture oligonucleotide #29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DM.
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                                                                                                                                                                                                 Hoppe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 4.2
); Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>ب</u>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Example; Page 5; 8pp; German.

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RESULT 908
ADKO1318/c
ID ADKO13
XX ADKO13
XX ADKO13
XX PAT DO -MAY
XX BS; hy
XX BS; hy
XX BS RATTUS
XX DE1020
XX DE008
XX DE068
XX DE68
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Comprise variable and non-variable regions. The capture oligonuclectides to have a 5' invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nuclectides to allow comprises are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, caids itvity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cailows imaging of the complete pattern of all nucleic acid in a cell, and can lows imaging of the complete spid, inexpensive and reproducible callows imaging of the comparison of nucleic acid pools, Since the method is based on comparison of nucleic acid pools. Since the capture probes used in the method of the hovention.
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Best Local Similarity
   Sorting single-stranded nucleic acid, useful for analyzing expression
                                                                        WPI; 2003-714082/68
                                                                                                                                             Boekenkamp D,
                                                                                                                                                                                                                                                                                    28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                  04-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rat DNA microarray capture oligonucleotide #38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK01318;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADK01318 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                                                                                                                                                                                                                                                                                                                                                         28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DE10208794-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          capture probes used in the method
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
                                                                                                                                                                                                                      DEGUSSA BIOACTIVES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            АЛАДАЛАЛАЛАБТАЛАЛ 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                             Dieck HT, Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.8; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of the invention.
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patterns and screening active agents, uses capture agent with variable and constant regions.

Example; Page 5; 8pp; German

This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and conjustives or supplements, especially minerals, trace elements, organic callows imaging of the complete pattern of all nucleic acid from cells. It can detect very small differences in pools of nucleic acid pool since the method is based on comparison of nucleic acid pools, not individual compounds in a cell, and capture probes used in the method of the invention.

Sequence 21 BP; 0 A; 1 C; 2 G; 18 T; 0 U; 0 Other;

Matches Query Match Best Local 16; h 1.18; Similarity 88.98; Conservative 0; Score 14.8; DB 1; Pred. No. 4.2e+02; Mismatches DB 1; Length 21; Indels 0; 0

밁 Ş 1520 AAAAAAAAAAAAGTAAAA 1537 18

06-MAY-2004 (first entry) ADK01323;

ADK01323 standard; DNA; 21

Rat DNA microarray capture oligonucleotide #43.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp

DE10208794-A1

04-SEP-2003

28-FEB-2002; 2002DE-01008794.

28-FEB-2002; 2002DE-01008794.

(DEGS) DEGUSSA BIOACTIVES GMBH.

PI XXX PA XXX PA

Boekenkamp D,

Dieck HT,

Hoppe H;

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RESULT 910
ADKO13
XX ADKO13
XX ADKO13
XX ADKO13
XX O6-MAY
XX 88; hy
XX 88; hy
XX Blood;
XX Blood;
XX BLOOD
XX BATCUB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25. T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC enducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single cranded nucleic acids to the surface is (quasi) covalent, supranolecular, CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpansive and reproducible cor allows imaging of the complete spide, inexpansive and reproducible of the complete pattern of all nucleic acid in a cell, and capture or permission of nucleic acid pool. Since the complete method is based on comparison of nucleic acid pools. Since the capture probes used in the method of the herention.
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Matches
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                                                                                                                                                                                                                                                                                                                                                               ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rat DNA microarray capture oligonucleotide #64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
   28-FEB-2002; 2002DE-01008794
                                                                    28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                           Rattue ep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADK01344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADK01344 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and constant regions.
                                                                                                                                                                                                                       DE10208794-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-714082/68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprise variable and non-variable regions. The capture oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DB 1; Length 21; Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gape
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RESULT 911
ADK01340/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC binding of particular sorts of single stranded nucleic acids. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The Capture CC capture oligonucleotides are biotinylated and immobilised on a surface by cinteraction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semicantestic, acids resided mucleic acids to the surface is (quasi) covalent, supramolecular, comparated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, continuity of pharmaceuticals and/or nutritional compounds, e.g. food cativity of pharmaceuticals and/or nutritional compounds, e.g. food cacids (amino, carboxylic or fatty acid) or their derivatives, salts and caids (amino, carboxylic or fatty acid) or their derivatives, salts and callows inaging of the complete sapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It callows inaging of the complete pattern of all nucleic acid in a cell, and can be based on comparison of nucleic acid pool. Since the method is based on comparison of nucleic acid pool. Since the method is based on comparison of sucleic acid pool. Since the method of the complete pattern of the nucleic acid pool. Since the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture
                                                                                                                                     ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                             Rat DNA microarray capture oligonucleotide #60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example; Page 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and constant regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                               06-MAY-2004
                                                                                                                                                                                                                                                                                                            ADK01340;
                                                                                                                                                                                                                                                                                                                                                        ADK01340 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1520 АЛААЛАЛАЛАЛАГТАЛАЛ 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21 AAAAAAAAAAAAAAA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                               (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.1%;
                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.8;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Migmatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.2e+02;
-has 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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04-SEP-2003 DE10208794-A1

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RESULT 912
ADRO1319/c
ID ADRO13
XX ADRO13
AC ADRO13
XX DT 06-MAY
XX DE Rat DN
XX
DE Rat DN
XX B9; hy
KW blood;
XX
XX Rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Compiles variable and non-variable region, the complement of which is present at CC have a 5'-invariable anchor region, the complement of which is present at CC that comprises all possible combinations of up to 10 nucleotides to allow CC binding of particular sorts of single stranded nucleic acids. The capture CC binding of particularly locked nucleic acids (LNA) and the anchor region CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The material and/or membrane, having semicant resin, gel, crystalline material and/or membrane, having semicant stranded nucleic acids to the surface is (quasi) covalent, supramolecular, comparison of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface by the method is used (i) for analysis of patterns, especially in mucosal, comparison, holod, nerve or germ cells and (ii) for determining the carids trone or supplements, especially minerals, trace elements, organic acids (amino, carboxyllc or fatty acid) or their derivatives, salts and carids (amino, carboxyllc or fatty acid) or their derivatives, salts and carids (amino, carboxyllc or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpensive and reproducible cor allows imaging of the complete spattern of all nucleic acid in a cell, and capture or permit of the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual capture prohes used in the method of the invention.
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       Rattus sp
                                               ss; hybridisation;
blood; nerve; germ
                                                                                                                  Rat DNA microarray capture oligonucleotide #39.
                                                                                                                                                                  06-MAY-2004
                                                                                                                                                                                                                ADK01319;
                                                                                                                                                                                                                                                             ADK01319 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 6; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Boekenkamp D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                              AAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                  (first entry)
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                                               capture oligonucleotide; pattern; mucosal; hair root;
cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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RESULT 913 ADK01328/c

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Matches Query Match Best Local (

16;

Conservative

0,

Similarity

1.1%;

Score 14.8; DB 1; Pred. No. 4.2e+02; Mismatches

DB 1; Length 21; Indels

<u>.</u>

Gaps

0

SEXEXEXE

ADK01328;

ADK01328 standard; DNA;

21 ВP

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #48.

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CC comprise variable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture objectives are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The Capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC siparticularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. CC The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the acids (anino, carboxylic or fatty acid) or their determining the cacids (anino, carboxylic or fatty acid) or their derivatives, salts and can detect very small differences in pools of nucleic acids from cells. It can detect very small differences in pools of nucleic acids from cells. It can detect very small differences in the nucleic acids from cell, and complete pattern of all nucleic acid in a cell, and complete pattern of all nucleic acid from cell, and complete pattern of solutional expresent comparison of nucleic acid pools. Since the complete status of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
Sequence 21 BP; 0 A; 2 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acids by isolation and hybridisation of nucleic acid pool reading out, where the nucleic acids are selectively bound using agents that are (a) immobilised on the surface of a solid matrix
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-714082/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Boekenkamp D,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example; Page 5; 8pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dieck HT, Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              describes a novel method for sorting single-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        poole,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             capture and (b)
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RESULT 914
ADK01335/c
ID ADK013
XX
AC ADK013
XX
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                                                                                                                                                                                                                                                           Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Boekenkamp D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-2002; 2002DE-01008794.
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blood; nerve; germ cell; food additive; food supplement.
                                                             ADK01335 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                              capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                    1520 ААААААААААААСТАААА 1537
                                                                                                                                                                18
                                                                                                                                                                                                                                           l Similarity
                                                                                                                                                                AAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                         BP; 0 A; 1 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                           1.1%;
                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                             Score 14.8; DB 1; Length 21; Pred. No. 4.2e+02;
                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                             Gaps
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ADK01335

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creading out, where the nucleic acids are selectively bound using capture cagnite that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable and non-variable regions. The capture oligonucleotides chave a 5'-invariable anchor region, the complement of which is present at the least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particulary sorts of single stranded nucleic acids. The capture capture oligonucleotides are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25. T residues. The capture oligonucleotides are biorinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, having the activity of pharmaceuticals and/or nutritional compounds, e.g. food cadditives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible capture probes used in the method of the invention. According from cells. It genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent conductive probes used in the method of the invention.
                                                                                                          Query Match
Best Local :
                                                                        Matches
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                                                                                                                                                                                                                     Sequence 21
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1520 AAAAAAAAAAAGTAAAA 1537
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                                                                                                              Similarity
                                                                    Conservative
                                                                                                                                                                                                                     BP; 0 A; 1 C; 1 G; 19 T; 0 U; 0 Other;
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                                                                        0
                                                                                                          Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                               Mismatches
                                                                                                                                                Length 21;
                                                                            Indels
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Rat DNA microarray capture oligonucleotide #22

06-MAY-2004 (first entry

ADK01302 standard; DNA; 21

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                                                                                                                                                                                        CY This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then CC reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides CC have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region CC agents are particularly locked nucleic acids (LNA) and the anchor region CC capture oligonucleotides are biotinylated and immobilised on a surface by conducting properties and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface stranded nucleic acids to the surface is (quasi) covalent, supramolecular, cc physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, that root, blood, nerve or germ cells and (i) for electrical sied or through a molecular sieve. The method provides rapid, inexpensive and reproducible caids (anino, carboxylic or fatty acid) or their derivatives, salts and cc andutives. The method provides rapid, inexpensive and reproducible the method is based on comparison of nucleic acid pools, not individual compones used in the method of the nucleic acid pool. Since the capture probes used in the method of the hypertion.
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Best Local S
Matches 16
                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
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                                1520 АДАЛАЛАЛАЛАД 1537
                                                                                      Similarity
                                                                                                                                                                                probes used in the
ANNANANANAAAAAAAA 1
                                                                                                                                              21 BP; 0 A; 0 C; 3 G; 18 T; 0 U; 0 Other;
                                                                       Conservative
                                                                                      88.9%;
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                                                                                                                                                                                    method
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                                                                                      Score 14.8; DB 1
Pred. No. 4.2e+02
                                                                       Mismatches
                                                                                                                                                                                    o
f
                                                                                                                                                                                    the invention.
                                                                                                           DB 1;
                                                                                                       Length 21;
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Matches Query Match Best Local :

Similarity

1.1%;

Score 14.8; DB 1; Pred. No. 4.2e+02; 0; Mismatches 2

DB 1; Length 21; Indels

0;

Gaps

0

Conservative

Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

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RESULT 916
ADKO1317/c
ID ADKO13
XX ADKO13
XX ADKO13
XX ADKO13
XX ADKO13
XX BS; hy
KW SS; hy
KW Blood;
XX Rattus
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                                                         cc nucleic acids by isolation and hybridisation of nucleic acids polls, then cc reading out, where the nucleic acids are selectively bound using capture capents that are (a) immobilised on the surface of a solid matrix and (b) cc comprise variable and non-variable regions. The capture oligonucleotides comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture capture oligonucleotides are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasilcovalent, supramolecular, cphysical, strimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the acids (amino, carboxylic or fatty soil or through a molecular sieve. The method provides rapid, inexpensive and reproducible captures. The method provides rapid, inexpensive and reproducible is allows imaging of the complete pattern of all nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acids from cells. It capture probes used in the method of the invention.

Cc apture probes used in the method of the invention.
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blood; nerve; germ cell; food additive; food supplement.
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                                                                                              This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is transported and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stransport and the surface is (quasi) covalent, supramplecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and carboxylic or fatty acid) or their derivatives, salts and carboxylic or fatty acid) or their derivatives, salts and carboxylic or fatty acid) or their derivatives, salts and carboxylic or fatty acid) or their derivatives, salts and mathod is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. Abx01281-Abx0
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probes used in the method of the invention

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CC comprise variable and non-variable regions. The capture oligonucleotides comprises variable anchor region, the complement of which is present at CC least comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture capture oligonucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The capture CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is sparticularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, cp hair root, blood, nerve or germ cells and (i) for through a molecular sieve. CC metal, resion, carboxylic or fatty acid) or their derivatives or supplements, especially minerals, trace elements, organic cardivitys of pharmaceuticals and/or nutritional compounds, e.g. food acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardivatives, salts and single field cardivatives, salts and cardivatives or supplements, especially minerals, trace elements, organic cardivity is method provides rapid, inexpensive and reproducible cardivity is magning of the complete pattern of all nucleic acids from cells. It callows imaging of the complete pattern of all nucleic acid in a cell, and calls based on comparison of nucleic acid pools, not individual
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cell; food additive; food supplement.
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                    This invention describes a novel method for sorting single-stranded CC reading out, where the nucleic acids are selectively bound using capture CC agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides CC have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture cagents are particularly locked nucleic acids (LNA) and the anchor region CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC conduction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, CC hair root, blood, nerve or germ cells and (ii) for determining the actids (amino, carboxylic or fatty acidly or their derivatives, salts and CC acids (amino, carboxylic or fatty acidly or their derivatives, salts and compounds, e.g. food
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Pred. No. 4.2e+02;
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This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture capture that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at comprise all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, cophysical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the
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germ cell; food additive; food supplement.
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Pred. No. 4.2e+02
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This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture capture state are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides chave a 5'-invariable anchor region, the complement of which is present at comparise once in each nucleic acid and a 3'-variable, discriminatory region contait comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are bictinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single
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Best Local :
This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (1MA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         physical, stimulated by an electrical field or through a molecular sieve The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, are can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                    Ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or through a molecular sieve.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            supramolecular,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hair root;
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metal, resin, gel, crystalline material and/or membrane, having semi-
conducting properties and especially in the form of a chip. Its surface
is particularly a layer of (bio)molecular filaments and binding of single
stranded nucleic acids to the surface is (quasi) covalent, supramolecular,
physical, stimulated by an electrical field or through a molecular sieve.
The method is used (i) for analysis of patterns, especially in mucosal,
hair root, blood, nerve or germ cells and (ii) for determining the
activity of pharmaceuticals and/or nutritional compounds, e.g. food
additives or supplements, especially minerals, trace elements, organic
acids (amino, carboxylic or fatty acid) or their derivatives, salts and
mixtures. The method provides rapid, inexpensive and reproducible
representation of differences in pools of nucleic acid in a cell, and
can detect very small differences in the nucleic acid in a cell, and
genes, matrix miniaturisation is possible. ADK01381-ADK01344 represent
capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid gools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilized on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; hybridisation;
blood; nerve; germ
                                                                                                                                                                                  Example;
                                                                                                                                                                                                                                                                                                                                          Boekenkamp D,
                                                                                                                                                                                                                                                                                                                                                                                                                        28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rat DNA microarray
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADK01304 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      interaction with streptavidin.
                                                                                                                                                                                                                       constant regions.
                                                                                                                                                                                                                                                                                                       2003-714082/68.
                                                                                                                                                                                                                                                                                                                                                                                 ) DEGUSSA BIOACTIVES GMBH.
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                                                                                                                                                                                                                                                              single-stranded
                                                                                                                                                                              Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 0 A; 1 C; 1 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                         screening active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                5
                                                                                                                                                                                                                                                                                                                                          Dieck HT,
                                                                                                                                                                              8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                capture oligonucleotide #24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.1%;
                                                                                                                                                                                                                                       nucleic acid, useful for analyzing expression active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                          Hoppe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The matrix is of plastic, ceramic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.2e+02;
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ADKO1306/c
ID ADKO13
XX
AC ADKO13
AC ADKO13
AX
XX Rat DN
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XX Blood;
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XX Blood;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                       Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Boekenkamp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-2002; 2002DE-01008794
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     blood; nerve;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rat DNA microarray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADK01306 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (DEGS )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1520 ААААААААААААСТАААА 1537
                                                                                                                                                                                                                                                                                                                                                                                              2003-714082/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18 AAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                              regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tion; capture oligonucleotide; pattern; mucosal; hair germ cell; food additive; food supplement
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            capture oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP
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Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, reading out, where the nucleic acids are selectively bound using cag agents that are (a) immobilised on the surface of a solid matrix and

pools, then

Example;

Page 5; 8pp;

German.

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RESULT 925
ADXO1325/c
ID ADXO13
XX ADXO13
XX C ADXO13
XX S8; hy
XX S8; hy
XX B1000;
XX B21020
XX D21020
YX D21020
YX D21020
YX D28-PEB
XX C CBGS
XX CDEGS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC have a 5'-invariable anchor region, the complement of which is present at CC have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow CC binding of particulary locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The material and/or membrane, having semi-CC conducting properties and especially in the form of a chip. Its surface by crystalline material and/or membrane, having semi-CC estimated nucleic acids to the surface is (quasi) covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino comparison of nucleic acid pools, inc cells. It captures arobase on comparison of nucleic acid pools, salts and CC and the based on comparison of nucleic acid pools, so the the comparison of nucleic acid pools. Since the capture arobase used in the method of the haventon of the hardron of individual comparison of nucleic acid pools.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                             Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                         DE10208794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          blood;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ss; hybridisation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADK01325;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADK01325 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 0 A; 1 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                          Boekenkamp D,
                                                                                                                                                                                                                                                                            28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                     28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprise variable and non-variable regions. The capture oligonucleotides
                                                                                                                                              WPI; 2003-714082/68
                                                                                                                                                                                                                                  (DEGS ) DEGUSSA BIOACTIVES GMBH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nerve;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tion; capture oligonucleotide; pattern; mucosal; hair root;
germ cell; food additive; food supplement.
                                                                                                                                                                                          Dieck HT,
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Pred. No. 4.2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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Example; Page 5; 8pp; German

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable

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RESULT 926
ADK01339/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       reading out, where the nucleic acids are selectively bound using capture comprise variable and non-variable regions. The capture oligonucleotides comprise a 5'-invariable anchor region, the complement of which is present at that comprises all possible combinations of up to 10 nucleotides to binding of particular sorts of single stranded nucleic acids. The capture oligonucleotides are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The materia is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, chipsical, stimulated by an electrical field or through a molecular sleve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the captures or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acid pool. Since the method is based on comparison of nucleic acid pools. Since the capture probes used in the method of the horizon acid pools. ADX01344 represent construct probes used in the method of the horizon in the surface acid pools.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, reading out, where the nucleic acids are selectively bound using cap agents that are (a) immobilised on the surface of a solid matrix and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                          28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                         04-SEP-2003.
                                                                                                                                                                                                                                                                                              DE10208794-A1
                                                                                                                                                                                                                                                                                                                                        Rattus
                                                                                                                                                                                                                                                                                                                                                                           ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Rat DNA microarray capture oligonucleotide #59.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-MAY-2004 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADK01339 standard; DNA; 21 BP
                                                        WPI; 2003-714082/68
                                                                                             Boekenkamp D,
                                                                                                                                     (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                               28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16; Conservative
                                                                                                                                                                                                                                                                                                                                      ф
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAAAAAAAAAAAAAA 1
                                                                                             Dieck
                                                                                             Ħ,
                                                                                             норре н;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pools, then
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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constant regions.

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RESULT 927
ADK01343/c
ID ADK013
XX ADK013
XX AC ADK013
XX O6-MAY
XX B8; hy
XX B8; hy
XX B8; hy
XX B100d;
XX DE1020
XX DE1020
XX DE1020
XX 28-FEE
XX 28-FEE
XX 28-FEE
XX 1DEGS
XX B0eker
XX BOeker
XX BOEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          F^{\times}_{\infty} \otimes S^{\times}_{\infty} \otimes S^{\times
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then CC reading out, where the nucleic acids are selectively bound using capture CC agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, carivity of pharmaceuticals and/or nutritional compounds, e.g. food acids (amino, carboxylic or fatty acid) or their derivatives, salts and carids (amino, carboxylic or fatty acid) or their derivatives, salts and carids (amino of differences in pools of nucleic acid from cells. It can lows imaging of the complete spid, inexpensive and reproducible method is based on comparison of nucleic acid pools, not individual capture nyoke sential on the method of the hearth of the invention. ADKOLD81-ADKOLD84 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                  Boekenkamp D,
                                                                                                                                                                                                                                                                                                      28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DE10208794-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rat DNA microarray capture oligonucleotide #63.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADK01343 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 0 A; 2 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      capture probes used in the method of the invention.
                                                                                                                                                                                    (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                        Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ..
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Comprise variable and non-variable regions. The capture oligonucleotides comprise variable and non-variable regions. The capture oligonucleotides comprise variable and non-variable regions. The capture oligonucleotides comprise once in each nucleic acid and a 3'-variable, discriminatory region comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture cidentices are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, comprised payer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, comprised payer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, comprised payer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, comprised payer of (bio)molecular field or through a molecular sieve. Comprised payer of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible compounds in a cell, and can detect very small differences in pools of nucleic acids from cells. It can detect very small differences in the nucleic acids from cells. It can detect very small differences in the method pools, so conditividual compounds. Since the matrix ministuries and co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 6; 8pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-714082/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and constant regions
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8
밁
                                       Matches
                                                 Query Match
Best Local :
                   1520 AAAAAAAAAAAGTAAAA 1537
                                       16; Conservative
                                                  Similarity
                                                  88.9%;
                                      0; Mismatches
                                                 Score 14.8; DB 1
Pred. No. 4.2e+02
                                                            DB 1;
                                       ر
ا
                                                           Length 21;
                                       0
                                       Gaps
                                       0
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Sequence 21 BP; 0 A; 1 C; 0 G; 20 T; 0 U; 0 Other;

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RESULT 928
ADK01301/c
                                                                ADK01301 standard; DNA; 21
                                                   06-MAY-2004 (first entry)
                                                          ADK01301;
                                                                ВÞ
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ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rat DNA microarray capture oligonucleotide #21.

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04-SEP-2003
                                     Rattus
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28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794

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RESULT 929
ADKO1312/c
ID ADKO13
XX ADKO13
XX ADKO13
XX DC Rat DN
XX BS; hy
KW BS; hy
KW BS; hy
KW BS; hy
KW BOOd;
XX Rattus
XX Rattus
XX DE Rattus
XX DE Rattus
XX DE RATTUS
XX DE O4-SEF
XX D4-SEF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         र्
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC comprise variable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow comprises as sequence of 10-50, particularly 15-25, T residues. The capture of comprises a sequence of 10-50, particularly 15-25, T residues. The capture of comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is physical, stimulated by an electrical field or through a molecular sieve. CC The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the catids (amino, carboxylic or fatty acid) or through a molecular sieve. CC minutures. The method provides rapid, inexpensive and reproducible caids (amino, carboxylic or fatty acid) or thrir derivatives, salts and cmid elect very small differences in pools of nucleic acids from cells. It can detect very small differences in nucleic acids from cells. It can detect very small differences in pools of nucleic acids from cells. It can detect very small differences in pools, not individual compounds is not individual compounds. Since the mucleic acid pools, not individual capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local (
                                                                                                                                                                                                                                                                       06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                               ADK01312 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                                                                                                                                                                                                                           Rat DNA microarray capture oligonucleotide #32.
                                                                                                                                                                                                                                                                                                                      ADK01312;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 1 A; 0 C; 2
                           04-SEP-2003
                                                                     DE10208794-A1
                                                                                                                                                           ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (DEGS ) DEGUSSA BIOACTIVES GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1520 AAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             норре н;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>..</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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RESULT 930
ADK01326/c
ID ADK013
XX
AC ADK013
XX
DT 06-MAY
XX
EX Rat DN
XX
KW 88; hy
XX
KW 88; hy
XX
XX
SX
Rattue
XX

06-MAY-2004

(first entry)

ADK01326;

ADK01326 standard; DNA; 21 BP

Rattus sp

ss; hybridisation;

tion; capture oligonucleotide; pattern; mucosal; hair root; germ cell; food additive; food supplement.

nerve;

Rat DNA microarray capture oligonucleotide #46.

밁 ঠ

18

AAAAAAAAAAAAAAAAA 1

Matches Query Match Best Local

16;

Conservative

<u>.</u>

Similarity

1.1%;

Score 14.8; Pred. No. 4 Mismatches

4.2e+02;

0

Gaps

0

DB 1; Length 21; Indels

1520 АААААААААААСТАААА 1537

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CC comprise variable and non-variable regions. The capture oligonucleotides comprise variable anchor region, the complement of which is present at CC least conce in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by CC interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filments and binding of single CC is particularly a layer of (bio)molecular filments and binding of single CC is particularly a layer of phastical field or through a molecular, CC physical, stimulated by an electrical field or through a molecular, CC hair root, blood, nerve or germ cells and (ii) for determining the CC acitivity of pharmaceuticals and/or nutritional compounds, e.g. food CC additives or supplements, especially minerals, trace elements, organic CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids maging of the complete sapid, inexpensive and reproducible cardetect very small differences in the nucleic acid scid in a cell, and CC activity probasics and comparison of nucleic acid pool. Since the method is based on comparison of nucleic acid pool. Since the capture probasic and the provides and possible. ADX01381-ADX01344 represent contributions are probasically and the invention.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sorting single-stranded nucleic acid, useful for analyzing expression parterns and screening active agents, uses capture agent with variable
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                                                        capture probes used in the method of the invention.
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  21
BP; 0 A; 0 C; 1 G; 20 T; 0 U; 0 Other;
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RESULT 931
ADK01305/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Comprise variable and of variable, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The comprises a sequence of 10-50, particularly 15-25, T residues. The comprises a sequence of 10-50, particularly 15-25, T residues. The comprise and surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semicontal components and surface by the metal, resinulated by an electrical fill or through a molecular, glass, and in the surface is (quasi) covalent, supramolecular, the method is used (i) for analysis of patterns, especially in mucosal, the method is used (i) for analysis of patterns, especially in mucosal, haviny of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxyllc or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpensive and reproducible captures and method is based on complete pattern of all nucleic acid in a cell, and captures are accomplete pattern of all nucleic acid in a cell, and capture of the complete solution of nucleic acid pools. Since the method is based on comparison of nucleic acid pools. Since the method is based on comparison of nucleic acid pools, not individual capture of the invention.
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se; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
                                         Rat DNA microarray capture oligonucleotide #25.
                                                                                    06-MAY-2004 (first entry)
                                                                                                                                 ADK01305
                                                                                                                                                                           ADK01305 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          capture probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8;
Pred. No. 4.2
                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                 4.2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length 21;
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ADK01310/ ID ADKO XX AC ADKO XX DT 06-M

ADK01310;

ADK01310 standard; DNA; 21

ВÞ

06-MAY-2004

(first entry)

RESULT 932

밁 Ś

Matches Query Match

16;

Conservative

0

Local

Similarity

1.1%;

Score 14.8; DB 1 Pred. No. 4.2e+02 Mismatches

DB 1; N 7:

Length 21; Indels

0

Gaps

0

Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

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CC comprise variable and non-variable regions. The capture oligonucleotides comparise variable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow comprises as sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids. The surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is physical, stimulated by an electrical field or through a molecular sieve. CC The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the acids (amino, carboxylic or fatty acid) or their derivatives, salts and caid surface is representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid from cells. It can detect very small differences in the nucleic acid from cells. It can detect very small differences in the nucleic acid from cells. It can detect very small differences in the nucleic acid from cells and capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
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Rat DNA microarray capture oligonucleotide #30

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then compared are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides chave a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region cc that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acide (LNA) and the anchor region cc agents are particularly locked nucleic acide (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, cc metal, resin, gel, crystalline material and/or membrane, having semi-cc is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, cc physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cardistives or supplements, especially minerals, trace elements, organic acids (amino, cardoxylic or fatty acid) or their derivatives, salts and cc additives. The method provides rapid, inexpensive and reproducible and method is based on comparison of nucleic acid pools, incells. It can lows imaging of the complete pattern of all nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADKOLZB1-ADKOL344 represent
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                                                                       Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                    Sequence 21 BP; 0
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blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and constant regions.
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                                  1520 AAAAAAAAAAAAGTAAAA 1537
18
                                                                                                                                                                                           probes used in the
AAAAAAAAAAAAAAAA 1
                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8pp; German.
                                                                                                                                                    A; 0 C; 2 G; 19 T; 0 U; 0 Other;
                                                                                         1.1%;
                                                                                                                                                                                           method of the invention.
                                                                         <u>,</u>
                                                                       Score 14.8; DI
Pred. No. 4.2e
0; Mismatches
                                                                                         4.2e+02
                                                                                                                DB 1;
                                                                                                            Length 21;
                                                                           Indels
                                                                         0;
                                                                           Gaps
                                                                           0
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RESULT 933 ADK01336/c ID ADK01336 standard; DNA; 21 BP.

Ś 밁

19

1520 AAAAAAAAAAAAGTAAAA 1537

Matches Query Match Best Local

Similarity

1.1%;

Score 14.8; Pred.

DB 1; Length 21;

<u>,</u>

Mismatches No. 4.2e+02

0

Gaps

0

Sequence 21 BP;

0 A; 0 C; 1 G; 20 T; 0 U; 0 Other;

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creading out, where the nucleic acids are selectively bound using capture comprise variable and non-variable regions. The capture oligonucleotides comprise variable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorte of single stranded nucleic acids. The capture combination granticular sorte of single stranded nucleic acids. The capture comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, comparises and especially in the form of a chip. Its surface conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, charing the method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the acids (amino, carboxylic or fatty acid) or their derivatives, eslits and captures. The method provides rapid, innerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and capture is maging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the mucleic acid from cells. It genes, matrix miniaturisation is possible. Abk01281-Abk01344 represent computer probes used in the method of the invention.
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blood; nerve; germ cell; food additive; food supplement.
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RESULT 934
ADKO1342/c
ADKO1342/c
ADKO133
XX ADKO13
XX ADKO13
XX ADKO13
XX BAPEN
XX CEAMP
XX Examp
XX E
                                                                                                  Computies variable and non-variable regions. The capture oligonucleotides to have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region CC computies a sequence of 10-50, particularly 15-25. T residues. The capture CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly acids to the surface is (quasi) covalent, supramolecular, cc physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cardistives or supplements, especially minerals, trace elements, organic acids (additives or supplements, especially minerals, trace elements, organic cardistics. The method provides rapid, inexpensive and reproducible and comparison of the complete acid pools, since the method is based on comparison of nucleic acid pools, Since the method is based on comparison of nucleic acid pools. Since the capture norbes used in the method of the heart of the invention.
                                                                               capture probes used in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rat DNA microarray capture oligonucleotide #62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement.
                                                                                  оf
                                                                                  the invention.
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Sequence

21

BP; 0 A; 0 C; 1 G; 20 T; 0 U; 0 Other;

Local Similarity

88.9%;

Score 14.8; DB 1 Pred. No. 4.2e+02 Mismatches

DB 1; Length 21; Indels

<u>,</u>

Gaps

0

Sequence 21 BP; 0 A; 1 C; 1 G;

19 T; 0 U; 0 Other;

Conservative

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ADKO1308/c
ID ADKO13
XX
AC ADKO13
XX
AC ADKO13
XX
BE ACTOM
XX
BE ACTOM
XX
BE ACTOM
XX
AC ADKO13

                                                              This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acid pools, then complete that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides chave a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture capture oligonucleotides are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, comprised by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic cativity of pharmaceuticals and/or nutritional compounds, e.g. food cativity of patterns of fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible cations in maging of the complete pattern of all nucleic acid from cells. It can detect very small differences in pools of nucleic acid from cells, and can capture probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 5; 8pp; German.
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germ cell; food additive; food supplement.
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Score 14.8; DB 1;

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RESULT 936
ADXO1311/c
ID ADXO13
XX ADXO13
XX B8; hy
XX B8; hy
XX B8; hy
XX B100d;
XX DT 06-MAY
XX B100d;
XX DE1020
XX DE1020
XX DE1020
XX DE1020
XX DE1020
XX DE1020
XX B0ekei
XX DEGS
XX WPI;
XX Sortil
PT B0ekei
XX WPI;
XX Sortil
PT partei
CC This
CC Comptr
C
This invention describes a novel method for sorting single-stranded cc nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramplecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, that root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardiscally in material and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardiscally in the nucleic acid from cells. It allows imaging of the complete pattern of all nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes.
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blood; nerve; germ cell; food additive; food supplement.
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RESULT 937
ADK01321/c
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This invention describes a novel method for sorting single-stranded conclusic acids by isolation and hybridisation of nucleic acid gools, then compared out, where the nucleic acids are selectively bound using capture capture that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides comprises variable anchor region, the complement of which is present at comprise once in each nucleic acid and a 3'-variable, discriminatory region combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids. The capture oligonucleotides agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are bictinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular single physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardivatives, salts and cardivatives.
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Pred. No. 4.2e+02;
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                           CC This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then CC reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonuclectides have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, certail, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quash) covalent, supramolecular, charity of sused (i) for analysis of patterns, especially in mucosal, capture of particular sieve. The method is used (i) for analysis of patterns, especially in mucosal, capture of the particular sieve or germ cells and (ii) for determining the
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                       have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture capture oligonuclest are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rat DNA microarray
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           patterns and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1520 АЛЛАЛАЛАЛАЛАСТАЛАЛ 1537
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single-stranded nucleic acid,
s and screening active agents,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tion; capture oligonucleotide; pattern; mucosal; hair root;
germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry
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RESULT 940
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or mutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid mixtures shall differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent
                                                                                                                                                                               Enriching low abundance polynucleotides in a sample, useful for gene expression analysis, comprises exposing the sample to an enzymatically non-extendable nucleobase oligomer to block polymerase activity on high
                                                                                                                                                                                                                                                                                                                                                                                                  13-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human ATP5F1
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                                                                                                                                                                                                                                                                                                                                             09-MAY-2002; 2002US-00144179.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               isoform 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADM96310 standard; DNA;
                                                                                                                                                                      abundance species.
                                                                                                                                                                                                                                                                                                                   (SCHR/)
                                                                                                                                                                                                                                                                                                       (CHEN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human; H+ transporting; mitochondrial ATP synthase; subunit form 1; ATP5F1; reverse transcriptase; RT-PCR; primer.
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Pred. No. 4.
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The invention relates to a method of enriching a low abundance polynucleotide in a sample of polynucleotides comprising a low abundance and a high abundance polynucleotide. The method comprises exposing the sample to an enzymatically non-extendable nucleobase oligomer having a nucleobase sequence complementary to a sequence within the high abundance polynucleotide under conditions so that base pairing occurs, and subjecting the sample to conditions for polymerase extension. Preferably, the enzymatically non-extendable nucleobase oligomer does not have a ribose-containing oligomeric structure. It is a peptide nucleic acid (PNA) oligomer or is a modified nucleotide oligomer or internucleotide

(EPIG-) EPIGENESIS PHARM INC

1; Page 20; 43pp; English.

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coligomers. The modified nucleotide oligomer or internucleotide analogue coligomers is selected from locked nucleic acids (LNA), N3'-P5' (C oligomer is selected from locked nucleic acids (LNA), N3'-P5' (C oligomer is selected from locked nucleic acids (LNA), N3'-P5' (C oligomers, C1-C4 alkylphosphonate oligomers, phosphoramidates, beta-C2 oligomers, C1-C4 alkylphosphonate oligomers, phosphoramidates, beta-C3 oligomers, C1-C4 alkylphosphonate oligomers, phosphoramidates, beta-C4 oligomers. The C1-C4 alkylphosphonate oligomers are methyl coligomer is chimeric. The sample comprises more than one high abundance polynucleotides. The sample comprises more than one high abundance coligomer is chimeric. The sample comprises more than one high abundance polynucleotide. The sample comprises more than one high abundance comprises excend strand cDNA synthesis. The sample is exposed to the nucleobase oligomer during the first strand/or second strand cDNA synthesis. The sample is exposed to the comprises second strand cDNA synthesis. The sample is exposed to the comprises oligomer during the first and/or second strand cDNA synthesis. The sample ocomprises on amplification step, which is by colombrase chain reaction (PCR) or by in vitro transcription. The RNA is calculated that the comprises of the method also comprises labelling the amplified polynucleotides. The method also comprises labelling the amplification. The comprises of labelling is concomitant with or subsequent to amplification. The comprise of subundance complised is expresent sequence represents a reverse transcriptise (RT)-PCR primer which was used to amplify the human import precursor of subunit B of the H+ transporting, mitochondrial ATP synthase, subunit B, isoform 1 (ATP5F1) companies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 941
ABD25907
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Best Local
                                                                                                                                                                                                                                                             Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 analogue oligomer. The modified nucleotide oligomer is selected from 2'-modified and 3'-modified nucleotide oligomers. The 2'-modified and 3'-modified nucleotide oligomers are selected from 2'-O-alkyl modified nucleotide oligomers and 3'-alkyl modified nucleotide oligomers. The 2'-O-alkyl modified nucleotide oligomers.
                                       24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABD25907;
                                                                                    23-APR-2002; 2002WO-US013143.
                                                                                                                                                                             WO200285309-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AI654215-derived oligonucleotide SEQ ID 4919.
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Compensation of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.

Combe invention also describes a kit, that comprises: (a) a delivery compensation also describes a kit, that comprises: (a) a delivery compensation also describes a kit, that comprises: (a) a delivery compensation also describes a kit, that comprises: (a) a delivery compensation also describes (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, and compensation of the kit. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung cinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, comprises and as pulmonary vasoconstriction, respiratory complements of the associated composition, allergies, asthma, impeded respiration, respiratory comprises syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary cransplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system c.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel composition (a) a first accomprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergreducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted inucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                 Selective enrichment;
                                                                                                                                                                                       RT primer used in the synthesis
                                                                                                                                                                                                                                  06-MAY-2004
                                                                                                                                                                                                                                                                          ADJ88057
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 15; SEQ ID NO 4919; 763pp; English
      22-JAN-2004
                                            US2004014105-A1
                                                                                     Unidentified.
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                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                       AAAAAAAAAAAAAAAA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                (first entry)
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L, Shahabuddin
                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 0 C; 0 G; 0 T; 0 U; 1 Other;
                                                                                                                                                 gene expression;
                                                                                                                                                                                                                                                                                                                   2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8;
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                                                                                                                                                                                       of an artificial gene transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2e+02
                                                                                                                                                 RT; reverse transcriptase; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pabalan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              first active
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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RESULT 943
ADM07216/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local (
                                                                                                                                                                        (CHEN/) CHEN C.
(SCHR/) SCHROEDER B
(BRAN/) BRANDIS J.
(SCHR/) SCHROTH G.
                                                                                                                               Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to methods for the selective enrice low abundance polynucleotides. The invention is useful for analy leaversesion in a sample and creating cDNA libraries. The present expression transcriptage (RT) primer used in the synthesis of an extraction of the synthesis of an expression of the synthesis of th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Enriching low abundance polynucleotide relative to a high abundance polynucleotide in a sample, for analyzing gene expression and creating cDNA libraries, comprises blocking polymerase activity on high abundance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA second :
differential
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Double-stranded cDNA synthesis; cDNA first strand synthesis; cDNA second strand synthesis; RNA template; RNA amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       is reverse transcriptase (kartificial gene transcript.
                                                                                      WPI; 2004-340131/31.
                                                                                                                                                                                                                                                                                                                                 23-OCT-2002; 2002US-00278760
                                                                                                                                                                                                                                                                                                                                                                                                                       US2004081962-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADM07216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; SEQ ID NO 41; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-121562/12.
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                                                                                                                                                                                                                                                                                   23-OCT-2002; 2002US-00278760
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polynucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1520 AAAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer used
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                                                                                                                               Schroeder B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene
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                                                                                                                                                                                                                      В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1, Pred. No. 4.2e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    မှု
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  strand synthesis.
                                                                                                                                 Schroth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ģ
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                                                                                                                                 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the selective enrichment of is useful for analysing gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2
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Synthesizing double-stranded cDNA, by synthesizing a cDNA strand from Ritemplate, removing the template and synthesizing double-stranded cDNAs using the cDNA as template in the presence of processive DNA polymerase

random

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RESULT 944
ADP09287/c
ID ADP092
XX ADP092
XX ADP092
XX Extend
XX Extend
XX Becret
KW Single
XX Homo 8
XX Homo 8
XX 10-JUN
XX 25-NOV
PR 25-NOV
PR 24-JUL
XX Roth R
PI Roth R
XX Roth R
XX Identi
PT or abs
PX Identi
PT or abs
PX Exampl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC primer complementary to template, removing the template, synthesising couble-stranded cDNAs in a second reaction mixture comprising processive CC DNA polymerase, DNA ligase, first cDNA strand as template and random CC primers having a mixture of oligonucleotides having random DNA sequences. CC Also disclosed is a method for amplifying a population of RNA molecules CC to produce a pool of double-stranded cDNA molecules, and a kit for CC synthesising quantitative information about the genetic profile of nucleic acid in original RNA sample. The method of the invention is CC useful in differential gene expression assays for the analysis of CC useful in differential gene expression assays for the analysis of CC diseased and normal tissue and for large-scale correlation studies on CC sequences, mutations, variants or polymorphisms among samples. The method is efficient in synthesising improved cDNA molecules and effective in CC generating useful quantities of an amplified cRNA product that comprises CC a population of cRNA molecules in substantially the same relative molar cratio as the RNA or mRNA starting material. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            breast cancer; cytostatic; gene therapy; human; chromogranin B; secretogranin 1; SCG1; chromosome 20pter-p12; ss; PCR; primer; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a method for synthesising double-
stranded cDNA, by synthesising first cDNA strands in a first reaction
mixture comprising reverse transcriptase, RNA template, and first strand
                      Example 5; Page 103; 286pp; English
                                                       Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                        25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
                                                                                                                                                                                                                                                                                                                      25-NOV-2003; 2003WO-US037966
                                                                                                                                                                                                                                                                                                                                                                                                    WO2004047767-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Extend primer 82 used to genotype human chromogranin B polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-AUG-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADP09287 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; SEQ ID NO 2; 19pp; English
                                                                                                                                           WPI; 2004-441082/41.
                                                                                                                                                                                                                                                                                                                                                               10-JUN-2004.
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                                                                                                                                                                                                                     SEQUENOM INC
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                                                                                                                                                                            Nelson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 0 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                              Braun
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                                                                                                                                                                              >
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.8; DB 1; Length 21; Pred. No. 4.2e+02; O; Mismatches 2; Indels
                                                                                                                                                                                Kammerer SM, Reneland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP;
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202020202020
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Best Local S
Matches 16
                                  Matches
                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cyrostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene three current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human chromogranin B (CHGB;secretogranin 1;SCG1) DNA which is located at chromosomal position 20pter-p12.
                                                                                                                      The invention relates to a novel human nucleotide reducing enzyme 59.62 polypeptide and encoding polynucleotide. The polynucleotide, polypeptide and its antagonist are useful for treating e.g. embryo development teratogenesis, blood system disease, and growth development disturbance disease. The present sequence represents the human nucleotide reducing enzyme 59.62 cDNA isolating RT-PCR primer
                                                                                                                                                                                                                                                                    Human nucleotide reducing enzyme 59.62 polypeptide and its encoding polynucleotide, for treating e.g. embryo development teratogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human nucleotide reducing enzyme 59.62 cDNA isolating primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 9 A; 5 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                               Sequence 24
                                                                                                                                                                                                                                            Example
                                                                                                                                                                                                                                                                                                                        WPI; 2002-305607/35
                                                                                                                                                                                                                                                                                                                                                       Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUL-2000; 2000CN-00117037.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-JUL-2000; 2000CN-00117037
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL60935;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL60935 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                     (SHAN-) SHANGHAI BIODOOR GENE DEV CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1311 TTTATTTTCAGAGACAGA 1328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               system disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 TTTTTTGCAGAGACAGA 4
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                                                   Similarity
    TGTTTTGTTTTTTAATCA 1267
                                                                                                                                                                                                                                          Page 17 (disclosure); 34pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                reducing enzyme 59.62; embryo development;
                                                                                               BP; 6 A; 2 C; 2 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.1%;
                                                   88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human; RT-PCR; primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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Pred. No. 4.2e+02;
                                                                Score 14.8;
                                                   Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                    Mismatches
                                                   o
O
                                                   .6e+02;
                                                                DB 1; Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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                                  0
                                                                                                                                                                                     polypeptide
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                                  Gaps
                                    0;
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RESULT 946
AAT69640/c

ID AAT69640;

XX

AAT69640;

XX

AAT69640;

XX

AT 20-FEB-1998 (first entry)

XX

DE Telomerase Oligo-dT-Primer P3.

XX

Telomerase oligo-dT-Primer; d

XW

Telomerase oligo-dT-Primer; d

XW

Telomerase oligo-dT-Primer; d

XW

Telomerase oligo-dT-Primer; d

XW

DE19644302-A1.

XX

DE19644302-A1.

XX

DE19644302-A1.

XX

DE19644302-A1.

XX

ABOEF ) BOEHRINGER MANNHEIM GMBH

XX

PH

ABOEF ) BOEHRINGER MANNHEIM GMBH

XX

BOEF ) BOEHRINGER MANNHEIM GMBH

XX

PH

CR

CR

Measuring telomerase activity, u

BOEF objected to be extending substration of the compound of the telomerase activity and the compound of the co
RESULT 947
ADM16445/c
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Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Measuring telomerase activity, useful for tumour diagnosis and compound screening - by extending substrate primer, followed by amplification and immobilising product for detection.
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      17-JUN-2004
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Pred. No. 5.1e+02;
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AAQ75722 standard;

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Matches 14
                                                                                                                                             The invention relates to isolated RNAs comprising an intron RNA that is released in a cell, thus modulating the function of a target gene. Also disclosed is a DNA template for the isolated RNA, an expression vector comprising the DNA, and a composition comprising one or more agents that induce RNA-mediated modulation of the functions of two or more agents that genes in a cell, such as a mammalian cell. The isolated RNAs and compositions are useful for modulating the function of a target gene in a cell, e.g. to inhibit a cancer-related gene, potential viral gene, and microbe-related gene, and thus useful for treating and preventing diseases such as cancer and viral/microbial infections. The current sequence represents a potential poly-pyrimidine tract of the artificial
                                                                                                                                                                                                                                                                                                                                            New isolated RNAs comprising an intron RNA that is released in a cell, thus modulating the function of a target gene, useful for treating and preventing diseases such as cancer and viral/microbial infections.
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12-OCT-2002; 2002US-0418405P.
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                                                                                                          Sequence 19 BP; 0 A; 3 C; 0 G; 0 T; 13 U;
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RESULT 949
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                                                                                                                                                                    Synthetic
                                                                                                                                                                                            Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                  Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                        (NITE ) NIPPON TELEGRAPH & TELEPHONE
                                                                16-APR-1993;
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                                                                              A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. Th method can be used to analyse gene expression rapidly and easily
                                                          Sequence 21
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         DE10208794-A1
                         Rattus sp.
                                                   ss; hybridisation;
                                                                   Rat DNA microarray capture oligonucleotide #5.
                                                                                       06-MAY-2004
                                                                                                        ADK01285
                                                                                                                         ADK01285 standard; DNA; 21
                                                                                                                                                                                                                                          Sequence
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                                                                                       (first entry)
                                          germ
                                                                                                                                                                                                                                                                                                                                                                                                                        93JP-00112515.
                                                                                                                                                                                                                                                                                                                                                                                                                                          93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                         capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement.
                                                                                                                                                                                                               1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                       0;
                                                                                                                                                                                                               Score 14.6; DB 1;
Pred. No. 4.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21
                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                         DB 1; Length 21
                                                                                                                                                                                                        0;
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                                                                                                                                                                                                         0
ADK01295
ID ADK
XX
AC ADX
AC ADX
XX
DT 06-
XX
XX
XX
DE Rat
XX
KW Bs;
KW blc
XX
                                                                                                                                                    S
                                                                                                                                                                                                         RESULT 953
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Matches Best

1302 TCTATTTTTTTTTTCAGA 1322

TTTTTTTTTTTTTTAGA 21

Local

Similarity

81.0%;

Score 14.6; Pred. No. 4. Mismatches

4.6e+02 DB 1;

Length 21; Indels

<u>,,</u>

Gaps

Ö

Conservative

<u>,</u>

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rat DNA microarray capture oligonucleotide #15

06-MAY-2004 ADK01295;

(first

entry)

ADK01295 standard;

DNA;

21

ΒP

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bo) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, having the cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, insxpensive and reproducible can detect very small differences in pools of nucleic acid from cells. It can be comparison of nucleic acid pools, not individual comparate probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-SEP-2003
Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example; Page 4; 8pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (DEGS ) DEGUSSA BIOACTIVES GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2003-714082/68.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hoppe H;
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Rattus

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RESULT 954
AAQ75762
BXAXAXB
                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel method for sorting single-stranded creading out, where the nucleic acids are selectively bound using capture cagents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture capture oligonucleotides are birthylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filanents and binding of single stranded nucleic acids. The capture capture oligonucleotides are birthylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filanents and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical filed or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the cacids (amino, carboxylic or fatty acid) or their derivatives, salts and cardives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and can detect very small differences in pools of nucleic acid from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the mucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual capture probes used in the method of the invention
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                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
                                        04-AUG-1995
                                                                                                                      AAQ75762 standard;
                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example; Page 5; 8pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-FEB-2002; 2002DE-01008794
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                                                                                                                                                                                                                                              1246 TCTTTGTTTTGTTTTTTAATC 1266
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                                                                                                                                                                                                                                                                                                       17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                        21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                      (first entry)
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             screening active agents, uses capture agent with variable regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dieck HT,
                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                      1.0%;
                                                                                                                          22
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                                                                                                                                                                                                                                                                                                                          Score 14.6; DB 1; Length 21; Pred. No. 4.6e+02;
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                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                       0
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Reverse transcription primer used in cDNA analysis technique

by digestion

enzymes.

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RESULT 955
AAQ75634
ID AAQ756
XX AAQ756
XX AAQ756
XX AAQ756
XX Revers
XX AAAlys
XX AAAlys
XX AAAlys
XX JP0630
XX JP0630
XX JP0630
XX JP0630
XX I6-APR
XX I6-APR
XX I6-APR
XX I6-APR
XX AAAlys
PF 16-APR
XX AAAlys
PF 1A-APR
XX AAAlys
PA (NITE
XX AAAlys
PT AAAlys
PT AAAlys
PT AAAlys
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                 Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                        aggregate;
                                                                                                                                                                                                                                                                           Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                      Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                     04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                AAQ75634 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 1 A; 2 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 8; 11pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JP06303997-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Analysis; gene expression; reverse transcription; primer; cDNA;
                                                            WPI; 1995-018287/03.
                                                                                        (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                      16-APR-1993;
                                                                                                                                                    16-APR-1993;
                                                                                                                                                                                  01-NOV-1994.
                                                                                                                                                                                                                JP06303997-A
                                                                                                                                                                                                                                                                                                                                                                                   AAQ75634;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Analysis of cDNA and gene expression - by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1246 TCTTTGTTTTGTTTTTTAATC 1266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     al Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 TTTTTTTTTTTTTTTCATC 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                        93JP-00112515.
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                                                                                                                                                    93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.6; DB 1.
Pred. No. 4.6e+02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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RESULT 956
AAQ75682
ID AAQ756
XX AAQ756
XX AAQ756
XX AAQ756
XX AAAJ9
CX Revers
XX Analys
KW Apgreg
XX Synthe
XX JP0630
XX JP0630
XX JP0630
XX I6-APF
XX (NITE
XX WPI; 1
XX WPI; 1
XX Analys
PT Analys
PT Analys
PT Analys
PT Analys
CC A metlo
CC double
CC and un
CC trans
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                                                                                                       Query Match
Best Local S
Matches 17
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abelled reverse transcription primers (GBNESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                   A method for the analysis of cDNA comprises (a) preparing an aggregate (double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEO files A0705547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                     Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 7; 11pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Reverse transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ75682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ75682 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1246 TCTTTGTTTTTTTTAATC 1266
                                                      1246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17;
                                                                                                                                                                                                                                                                            can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       μ
                                                                                                          l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                  TCTTTGTTTTTGTTTTTTTTTTTAATC 1266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTTTTTTTTTTTTTTGATC
TTTTTTTTTTTTTTTATTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene expression; reverse transcription;
; restriction enzyme; ss.
                                                                                                                                                                                                                                                                         be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                     BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                 1.0%;
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Pred. No. 4.6e+02;
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Best Local :
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16-APR-1993;
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            A method for the analysis of cDNA comprises (a) preparing an aggregate (double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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17; Conservative
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                           TCTATTTTTTTTTTCAGA 1322
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.6;
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                                                                                                                                                                                                                                                                                                                                                                                                                 4.6e+02;
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Reverse transcription primer

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  Disclosure; Page 7; 11pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-018287/03.
                            Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                           16-APR-1993;
                                                                                                                                                    16-APR-1993;
                                                                                                                                                                                 01-NOV-1994
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                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                               aggregate;
                                                                                                                                                                                                                                                                            Analysis;
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                                                                                                                                                                                                                                                                                                                                  04-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Analysis of cDNA and gene expression – by amplification of mRNA followed by digestion with restriction enzymes.
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17; Conserv
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                                                                                                                           93JP-00112515
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Pred. No. 4.6e+02;
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                                                  The present invention describes a method for the comparative detection of the amount of an RNA. The method comprises: (a) cDNAs obtained by CC transcribing respectively from at least two tissue RNAs are respectively CC fragmented by using a same restriction enzyme; (b) each different adaptor are added to each of the cDNA fragments derived from CC the same or different tissues by the step (a); (c) the resultant adaptor added cDNAs are mixed together; (d) an adaptor primer having the common CC to amplify said adaptor-added cDNAs containing no region derived from CC polyadenylic acid of the mRNA before the addition of the adaptor among CC cDNA amounts are measured between the tissues; (f) the RNA is detected CC from the measured result; (g) each different adaptor and a common adaptor cadded to BNAs prepared by the step (b); (e) the ratios of the CC different individuals; (h) the resultant adaptor-added genomic DNAs are mixed together; (i) the adaptor-added genomic DNAs are mixed together; (i) the adaptor-added genomic DNAs are of the genomic DNAs are measured between the individuals. The method is common control to the defection of the amounts of the amplified amounts of the genomic DNAs are measured between the individuals. The method is common for the detection of the amounts of RNA and DNA. The present
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Comparative detection of the amounts of RNA and DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-MAY-2000; 2000JP-00160324
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                                     sequence represents an oligonucleotide
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Pred. No. 4
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                                     e individuals. The IA and DNA. The IA is used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4
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of the present invention

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RESULT 964
AA166361
ID AA1663
XX Human;
KW Human;
KW Haemop;
KW Inflam
XX Homo a
XX Ho
RESULT 965
ABK86169/c
ID ABK861
XX
AC ABK861
XX
DT 24-SEP
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides the protein and coding sequences of human phosphatidylinositol-3 (PTDINS-3) kinase 35. The sequences can be used in the treatment of cancer, haemopathy, HIV infection, development disorders, immunological diseases and inflammation. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New human phosphatidylinositol-3 (PTDINS3) kinase 35 for diagnosing and treating malignant tumor, hemopathy, human immunodeficiency virus infection, immunological diseases and various inflammations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; phosphatidylinositol-3 kinase 35; PTDINS-3 kinase 35; cancer; haemopathy; development disorder; HIV infection; immunological disease; inflammation; gene therapy; PCR primer; ss.
                            24-SEP-2002
                                                                                 ABK86169;
                                                                                                                                 ABK86169 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 12; 34pp; Chinese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-MAR-2000; 2000CN-00114973.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  is a PCR
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                                                                                                                                                                                                                                                                                                                                                           Local Similarity
168 17; Conserv
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer for the coding sequence of the invention
                                                                                                                                                                                                                                                                                                         TCTATTTTTTTTTTTCAGA 1322
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 3 A;
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                                                                                                                                                                                                                                                                                                                                                              Conservative
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                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                   1.0%;
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Pred. No. 4e
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
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RESULT 966
ABK86168
ID ABK861
XX
AC ABK861
XX
DT 24-SEP
XX
DE Oligo
XX

standard; DNA; 24 BP

24-SEP-2002 ABK86168 ABK86168

(first entry)

Oligo dT primer #1 used in

method

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study gene expression.

밁 S

21

TTTTTTTTTTTTTTAAAC

1246 TCTTTGTTTTTTTTTAATC 1266

Query Match Best Local Matches

al Similarity 81.0%; 17; Conservative

0 Pred. Score 14.6; Mismatches

No.

4e+02; DB 1; 4.

Length Indels

24;

0

Gaps

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The invention relates to systems for identification and characterisation of gene expression in one or more samples, comprising an identimer having a specific oligo-dT primer sequence, where the identimer comprises a condition of detectable marker at its 5° end. The system is useful for identifying any or all genes expressed in a given in vivo or in vitro RNA sample, as well as the relative differences in mRNA between 2 or more samples, where condesired, for supporting discovery of new genes, and for identifying mRNAs that are expressed at different levels between 2 or more samples. The new system or method addresses limitations of prior methods by comprising compositions and systems that incorporate new strategies where molecular or biochemical assay compositions and systems are linked to DNA or RNA sequence databases for optimal resource efficiency in assaying gene expression. The system has the following advantages over existing methods: (a) prior sequence information or clone library construction is not needed to enable the assay; (b) provides immediate sequence in formation in addition to information concerning changes or differences in one assay; (c) generates cDNA fragments from all mRNAs present in the sample for subsequent investigation by common molecular biology techniques; and (d) does not require prior knowledge of the sequence of the sequence of the organism under investigation and can be employed in organisms lacking significant genomic sequence in formation. The present convention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Identifying and characterizing gene expression in samples, identifying mRNAs expressed at different levels, comprises identimer having a oligo-dT primer of a specific sequence detectable marker at its 5' end.
Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-508123/54.
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BP; 20 A; 0 C; 1 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        45pp; English.
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RESULT 967
ADB04575/c
ID ADB045
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AC ADB045
XZ
DT 20-NOV
XX
DE Human
XX
KW Cytost
KW zinc f
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to systems for identification and characterisation CC of gene expression in one or more samples, comprising an identimer having CC a specific oligo-dT primer sequence, where the identimer comprises a cC detectable marker at its 5 end. The system is useful for identifying any CC or all genes expressed in a given in vivo or in vitro RNA sample, as well cas the relative differences in mRNA between 2 or more samples, where CC desired, for supporting discovery of new genes, and for identifying mRNAs cC that are expressed at different levels between 2 or more samples. The new system or method addresses limitations of prior methods by comprising CC compositions and systems that incorporate new strategies where molecular CC or biochemical assay compositions and systems are linked to DNA or RNA CC sequence databases for optimal resource efficiency in assaying gene CC expression. The system has the following advantages over existing CC methods: (a) prior sequence information or clone library construction is information in addition to information concerning changes or differences CC in one assay; (c) generates CDNA fragments from all mRNAs present in the cample for subsequent investigation by common molecular biology ct techniques; and (d) does not require prior knowledge of the sequence of the organism under investigation and can be employed in concernion represents an oligo dT primer used in the method of the
                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                      Best Local Similarity Matches 17; Conserv
 Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying and characterizing gene expression in samples, for identifying mRNAs expressed at different levels, comprises employing an identifier having a oligo-dT primer of a specific sequence and a detectable marker at its 5' end.
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                                                       Human MDZ7 scanning oligonucleotide SEQ ID 5561.
                                                                                             20-NOV-2003
                                                                                                                                  ADB04575
                                                                                                                                                                    ADB04575
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 24 BP; 3 A; 1 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                      standard; DNA; 25
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                                                                                                                                                                                                                                                                                                                                    1.0%;
nilarity 81.0%;
Conservative
                                                                                             (first entry)
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                         Score 14.6;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                       4e+02;
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                                                                                                                                                                                                                                                                                                                                                                        Length 24;
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AAN50186/c
 RESULT 968
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer; developmental disorder; ss.
                                                                                                                                                                                                25-MAR-2003
01-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 25 BP; 4 A; 2 C; 3 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shannon M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JUL-2002; 2002EP-00016874.
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06-NOV-1984;
                                  03-JUL-1985
                                                                EP146743-A.
                                                                                               Homo sapiens
                                                                                                                            Immunoglobulin
                                                                                                                                                              Sequence of human immunoglobulin gene enhancer region called unit
                                                                                                                                                                                                                                                AAN50186;
                                                                                                                                                                                                                                                                                AAN50186 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 8; SEQ ID NO 5561; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                (revised)
(first entry)
84EP-00113371
                                                                                                                               gene; enhancer; promoter; expression cassette;
                                                                                                                                                                                                                                                                                DNA;
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81.0%; Pred. No. 3.
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06-DEC-1983;
04-JUN-1984;
05-JUN-1984;
                                                                                                                                                                                                                                                         KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; tumour angiogenesis; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The inventors claim a gene fragment comprising (a) the enhancer DNA segment and (b) a structural gene such as human D,V and J gene, and a promoter. Using the gene fragments of the invention, transcription and translation efficiency are enhanced. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kudo
                                stability
                                                             WPI, 1997-259017/23.
                                                                                                                                   26-OCT-1995;
11-JAN-1996;
                                                                                                                                                                 25-OCT-1996;
                                                                                                                                                                                                          WO9715662-A2
                                                                                                                                                                                                                                                                                           Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1;
                                                                                                                                                                                                                                                                                                                Human flt1 VEGF receptor hammerhead ribozyme substrate #1090.
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                                                                                                                                                                                                                              Homo sapiens
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15; Conserv
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CHIRON CORP
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                                         acid molecule modulating VEGF
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    useful

                                                                                Mcswiggen J,
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                                                                                                                                                                                                                                                  kinase 1; ss.
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83JP-00229037.
84JP-00113047.
84JP-00113834.
                                                                                                                                   95US-0005974P.
96US-00584040.
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                                                                                                              PHARM INC
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93.8%;
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Pred. No. 6.6e+02
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                               F receptor(s)
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                              angiogenesis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Watanabe
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                                        gene expression
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Claim 4;

Page

79; 218pp; English

rheumatoid arthritis, etc.,

in a human

patient

to AAA19086

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ARASILIT 970
ARASIJ31/C
ID ARASIJ3
AC ID ARASIJ3
AC ARASIJ3
AC ID ARASIJA
AC ID ARAS
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Best Local S
Matches 15
The present invention describes enzymatic nucleic acid molecules with RNJ cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, and AAA17168 to AAA17560 and AAA17633 to AAA17644 represent their corresponding target sequences; AAA17685 to AAA13836 and AAA1987 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA1908 and AAA19155 to AAA19222 represent their corresponding target sequences;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes nucleic acid molecules which modulate t synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antifilammatory; antiarthritic; antisoriatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-OCT-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-591315/50
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                                                                                                                                                                                                                                                                                                                                                                             ribozymes for modulating the synthesis, mRNA encoding an angiogenic factors.
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                                                                                                                                                                                                                                                                                                                  Page 200; 305pp; English.
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Pred. No. 6.2e+02;
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Best Local
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Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; 8s.
                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-OCT-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Integrin alpha 6 subunit substrate sequence SEQ ID NO:4538.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-JUN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA21312 standard; RNA; 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1512
                                                                                                                                                                                                                                                                                                               1999-591315/50
                                                                                                                                                                                                                                                                                                                                                                            PA,
                                                                                                                                                                                                                mRNA encoding an angiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17
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                                                                                                                                                                                                                                                                                                                                                                                                                                        RIBOZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGGTAATTAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TGTTAATTAAAAAAAA 1527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             subunit alpha-6, or integrin subunit beta-3
                                                                                                                                                                                                                                                                                                                                                                        Roberts E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                       200; 305pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98US-0079678P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-US006507
                                                                                                                                                                                                                                                                                                                                                                                                                                        PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                        Jarvis T,
                                                                                                                                                       English
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Pred. No. 6
                                                                                                                                                                                                                                                   the synthesis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                            Coeshott
                                                                                                                                                                                                                       factors
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                                                                                                                                                                                                                                                   expression and/or stability
                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
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The present invention describes enzymatic nucleic acid molecules with cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to

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AAV91364
ID AAVY
XX
AC AAVS
XX AAVS
XX Humm
DE Humm
XX Humm
KW Humm
KW Infi:
KW rest
XX Homm
XX WO9
XX WO9
XX O9-1
PR 09-1
PR 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, CC and AAA17168 to AAA17560 and AAA17623 to AAA17684 represent their CC corresponding target sequences; AAA17685 to AAA18385 and AAA19087 to CC AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 CC AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA19086 CC AAA19155 to AAA20361 and AAA21501 to AAA21595 represent ribozyme sequences; CC AAA19222 represent their corresponding target sequences; CC AAA21596 to AAA21688 represent their corresponding target sequences; CC AAA21596 to AAA21688 represent their corresponding target sequences; CC AAA21596 to AAA22475 and AAA22363 to AAA2362, AAA22343 to CC AAA21689 to AAA22475 and AAA22476 to AAA2362, AAA23343 to CC AAA23422 represent their corresponding target sequences. The ribozymes of CC the invention are used for modulating the synthesis, expression and/or CC the invention are used for modulating the synthesis, expression and/or CC their subunit beta-3, integrin subunit alpha-6, or Tie-2. They are cspecially used to treat cancer, diabetic retinopathy, age related CC macular degeneration (ARMD), inflammation, and arthritis, as well as CC angiofibroma of tuberous sclerosis, pot-wine stains, Sturge Weber Syndrome, Kippel-Treanunay-Weber Syndrome, Osler-Weber-Rendu syndrome, CC and other syndromes and diseases related to the levels of ARNT, Tie-2, integrin subunit alpha-6, or integrin subunit beta-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                         Jarvis T, M
Parry T, Be
Thompson J,
                                                                                                                                                                                                                                                                                 02-OCT-1997;
02-OCT-1997;
05-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                           09-MAY-1997;
09-JUN-1997;
03-JUL-1997;
22-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification, synthesis; deprotection; purification; cancer; inflammation; psoriasis; non-hepatic ascites; infection; genetic drift;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human C-raf target site nucleotide position 2745.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 restenosis; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV91364 standard; RNA; 17
                                                                                                                                                                                          (RIBO-)
                                                                                                                                                                                                                                                        19-DEC-1997;
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                                                                                                                                                                                          RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 93.8
15; Conservative
                                                         Maturra L, Ma
Beigelman L, Ma
Morkman CT,
                                                             Matulic-Adamic J, Reynolds M, Kisich K,
Beigelman L, Mcswiggen JA, Karpeisky A,
, Workman CT, Beaudry A, Sweedler D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 4 A; 2 C; 0 G; 0 T;
                                                                                                                                                                                                                                                 97US-0046059P.
97US-0049002P.
97US-0051718P.
97US-0056808P.
97US-0061321P.
97US-00643266P.
97US-0068212P.
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No. 6.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
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ARESULT 973
ARZ38362/c
ID ARZ383
XX ARZ383
AC 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SP1; consensus; competition gel mobility shift assay; apolipoprotein apo AI; low density lipoprotein; LDL; cholesterol; coronary artery disease; promter; flanking region; plasmid; DRE; drug responsive element; reporter gene; drug screening; candidate; expression; identification; rapid; automation; enhancer; element; ds.
Screening for drugs that increase expression of the apolipoprotein AI gene, which may then be useful for treating coronary artery disease.
                                                                                                                                                                                                                                                                                                                                   29-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                               WPI; 2000-038251/03
                                                                                                                                                                                                                                                                                                                                                                                                   29-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US5994061-A
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RESULT 974
AAA25444/c
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Best Local (
                                                                                                                                                                                                                                                                                                                                   Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17
                                                Matulic-Adamic
                                                              Thompson JD, Beigelman L, Reynolds M, Zwick M, Jar
                                                                                                                                                20-APR-1998;
23-JUN-1998;
                                                                                                                                                                                                                                                                                                     Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA25444 standard; DNA; 17
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                                                                                                                 (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                   19-APR-1999;
                                                                                                                                                                                                                                    28-OCT-1999.
                                                                                                                                                                                                                                                                    WO9954459-A2
                                                                                                                                                                                                                                                                                                   sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                     receptor hammerhead ribozyme target sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 2 A; 4 C; 10 G; 1 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                98US-0082404P.
98US-00103636.
                                                                                                                                                                                                   99WO-US008547
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                                                                n L, Mcswiggen JA,
Jarvis T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Haeberli
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WPI; 2000-013248/01

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 975
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Best Local
                                                                                                                                                       Thompson JD, Beigelman L, Mcswiggen JA, Reynolds M, Zwick M, Jarvis T, Woolf T, Matulic-Adamic J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease;
                                 New nucleic acids that
                                                                                                   WPI; 2000-013248/01.
                                                                                                                                                                                                                                                                                                                                                       20-APR-1998;
23-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1944
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acids that interact, and optionally cleave, target sequences,
                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1521 AAAAAAAAAAAGTAAA 1536
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15; Conserv
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98US-00103636.
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                                    interact,
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Pred. No. 6.2e
0; Mismatches
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                                    and optionally cleave, target sequences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4;
. 6.2e+02;
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Haeberli
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RESULT 976
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Best Local
                                                                                                                        Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor prointerferon alpha and erythropoietin.
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                                                                                                                                                                                                                       WPI; 2000-647423/62.
                                                                                                                                                                                                                                                                 Blatt L,
                                                                                                                                                                                                                                                                                                                                                         12-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ribozyme; erythropoietin; granulocyte colony stimulating factor;
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                                                                                   Claim 42; Page 123; 164pp; English.
                                                                                                                                                                                                                                                                                                               (RIBO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   interferon alpha;
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                                                                                                                                                                                                                                                                 Pavco P,
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Pred. No. 6.2e+02
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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription

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RESULT 977
AAFO6140
ID AAFO61
XX AAFO61
XX AAFO61
XX Hammer
XX Ribozy
KW Ribozy
KW Ribozy
KW Interf
XX Homo a
XX Claim
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Blatt L,
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                                                                                                                                                                                                      Sequence
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                                              CTTTGTTTTGTTTTTT 1262
силивиливилении 17
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RESULT 978

Sequence 17

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                                                                                The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 concleic acids can be used as probes to detect, characterise and quantify hgdMLP-1 nucleic acids in samples, as amplification substrates, to comprove the initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for capture sing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as thardards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as tharpseutic supplement in pattients having specific deficiency in hGDMLP-1 compounds a consentration, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a conjuncted associated with the expression of hGDMLP-1, in particular heart can skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

The present sequence in the exemplification of the present invention. N.B. the sequence data for this patent did not form part of the printed capture data for this patent did not form at directly from WIPO are fermined.
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30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
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30-JAN-2001;
30-JAN-2001;
The present invention describes a human genome-derived myosin-like protein 1 (hgDMLP-1). The protein and polynuclectide sequences of hgDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1
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1467/c
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nilarity 93.8%;
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2000US-023458PP.
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2001WO-US000661.
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Pred. No. 6.2e+02;
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ARBNO7279/C
ID 28N0722
XX ABN0722
XX 29-MAY
XX Human;
XX Human;
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XX M Human;
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30-JAN-2001;
30-JAN-2001;
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21-SEP-2000;
27-SEP-2000;
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30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000666.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
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                                                                                                                    New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine;
                                                                         Disclosure; SEQ
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30-JAN-2001;
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                                                                                                                                                                                                                                WPI; 2002-179446/23.
                                                                                                                                                                                                                                                                                                                                    (AEOM-)
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; 2001WO-US000670.
; 2001US-0266860P.
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2001WO-US000663.
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2000US-0236359P.
2000GB-00024263.
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                                                                       ID NO 2271; 214pp;
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Pred. No. 6
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                                                                         English.
                                                                                                                                                                                                                                                                               Rank DR,
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                                                                                                                                                                                                                                                                                  Shannon
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The present protein 1 (

1 (hGDMLP-1).

invention describes

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protein and po

an genome-derived myosin-like polynucleotide sequences of l

hGDMLP-

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CC 1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify CC hGDMLP-1 nucleic acids in samples, as amplification substrates, to CC provide initial substrates for the recombinant engineering of hGDMLP-1 CC expressing the proteins. The hGDMLP-1 proteins or polypeptides may be CC used as immunogens to raise antibodies that specifically recognise hGDMLP-1 concentration CC and/or amount specifically of hGDMLP-1 proteins, as specific biomolecule CC capture probes for surface-enhanced laser desorption ionisation, as CC therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The CC polynuclectide sequences encoding hGDMLP-1 may be used for diagnosing a CC disorder associated with the expression of hGDMLP-1, in particular heart CC and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. CT he present sequence represents an ollower used in the screening of the present sequence in the expression of the present invention. N.B. CT he sequence data for this patent did not form part of the printed CC specification, but was obtained in electronic format directly from WIPO CC at fifth the present sequence with the screening of the specification, but was obtained in electronic format directly from WIPO CC at fifth the printed CC specification, but was obtained in electronic format directly from WIPO CC at fifth the printed content the present invention.
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Best Local :
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21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                                                                                                                                                                                     30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; genome-derived myosin-like protein 1; GDMLP-1; hGDMLP-1; heart; muscle; myosin; chromosome 22; gene therapy; vaccine; heart disease; skeletal muscle disorder; amplicon; screening; ss.
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                                                                                                                                                                                                                                                 30-JAN-2001;
30-JAN-2001;
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                                            (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                           30-JAN-2001;
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    Ji Y,
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                                                                                 2001WO-US000664.

2001WO-US000665.

2001WO-US000666.

2001WO-US000667.

2001WO-US000668.

2001WO-US000669.

2001WO-US000670.

2001US-0266860P.
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2001WO-US000661.
2001WO-US000662.
2001WO-US000663.
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2000US-0234687P.
2000US-0236359P.
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    Penn SG,
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  Hanzel
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    Rank DR,
    Chen
    Ξ
    Shannon
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                                                                                                                                                                                                                    therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp. wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                              and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
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                                          396 GCCGAAGGCCCGCAGG 411
17 GCCGAAGGCCCGGAGG
                                                                                        15; Conservative
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Pred. No. 6.2e+02;
0; Mismatches 1;
                                                                                                                                                                                     0 Other;
                                                                                                                                   Length 17;
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                                                                                          <u>,,</u>
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RESULT 982
ACN12480
(RIBO-) RIBOZYME PHARM INC.
(BLAT/) BLATT L.
(MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                                                                                     MNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WNV minus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACN12480 standard; RNA; 17
                                                                                                                                                     19-OCT-2001; 2001WO-US048350.
                                                                                                                                                                                                          06-SEP-2002
                                                                                                                                                                                                                                                                                                          West Nile Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACN12480;
                                                                                                     20-OCT-2000; 2000US-0242411P.
                                                                                                                                                                                                                                                                                                                                                             Amberzyme;
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                                                                                                                                                                                                                                                                                                                                                             Zinzyme;
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RESULT 99
ACNOLLIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to MNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nucleicides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virucide; neuroprotective; antibacterial encephalitis; myocarditis; meningitis; iliver failure; cancer; cirrhosis; Hammer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis;
                                                                                                                      WPI; 2002-706994/76.
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(BLAT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           West Nile Virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WNV Hammerhead Ribozyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-APR-2004
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                                                                                                                                                                                                                                                                         RIBOZYME PHARM INC
BLATT L.
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                                                                                                                                                                                                                                                MCSWIGGEN J A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCAGGAUGUAACCCAG 17
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                                                                                                                                                                              Mcswiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zinzyme;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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Pred. No. 6.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myotarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                     New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                                                                                                                                                           WPI; 2002-706994/76.
                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J A.
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                                                                                    1199; 495pp;
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Pred. No. 6.2e+02
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The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (MNV). The nucleic acid molecules are useful for treating a condition related to MNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis,

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The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to wNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocallular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080
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                                                                                                                                                                                                                                                                                                                                                                                                                        New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L.
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                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 14423; 495pp; English.
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Pred. No. 6.2e-
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Query Match

Score 14.4;

DB 1;

Length 17;

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Matches 15
                         The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (MVV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, mycarditis, meningitis, neurologic infection, hepatitis, niver failure, hepatocellular carcinoma or cirrhosss. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic molety. Nucleic acid molecules SEQ ID NO 1194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid molecule of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           virucide; neuroprotective; antibacterial; replication; pancreatit encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme; Amberzyme; Zinzyme; ss.
                                                                                                                                                                                                                                                              New nucleic acid molecule that modulates replication of West Nile Vi (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-)
(BLAT/)
  Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            West Nile Virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               are claimed; however, in the specification. molecule of the invent
                                                                                                                                                                                                                                                                                                                                                 Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                20-OCT-2000; 2000US-0242411P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WNV Inozyme substrate SEQ ID NO 3019.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-APR-2004
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BLATT L.
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                                                                                                                                                                                                                                    SEQ ID NO 3019; 495pp; English.
                                                                                                                                                                                                                                                                                                                                                 Mcswiggen
  BP; 3 A;
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Pred. No. 6.2e+02
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    0 T; 4
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                                                             Matches
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                                                                                                                                                                 of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least ten 2'-O-methyl modifications, phosphorothioate linkages on at least three of the 5' terminal nucleotides and a 3' end modification of a 3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                  New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                       Sequence 17
                                                                                                                                                                                                                                                                                                                                                           The invention relates to nucleic acid molecules that modulate replication
                                                                                                                                                                                                                                                                                                                                                                                          Claim 23; SEQ ID NO 7862; 495pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-706994/76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-APR-2004
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(BLAT/) BLATT L.
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                              CGCTGGTGAATGGAGT 711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mcswiggen JA;
                                                                                                                        BP; 4 A;
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                                                                           1.0%;
                                                                                                                        7 C; 3 G; 0 T;
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                                                                          Score 14.4; DB 1;
Pred. No. 6.2e+02;
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                                                                                                                        3 U; 0 Other;
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ARESULT 988
ARESULT 988
ARE 360360
AC ARE 36
                                                                                                                                                             The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive CC alignment, at least 80 % identity to the 17 mer sequence with, after optimal CC alignment, at least 80 % identity to the 17 mer sequence that CC of any of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, CC identifying, quantifying and/or amplifying a nucleic acid, e.g. as one CC component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, CC polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for CC preparation of pharmaceuticals for prevention and/or treatment of viral CC diseases that are characterised by development of tumours or cell CC diseases that are characterised by development of tumours or cell CC diseases. The polypeptides can also Alabeimer's disease and CC schizophrenia. Analysis of the expression of the 17 mer nucleic acids in gatient samples is useful for diagnosis and/or prognosis of these CC both the polypeptide and antibodies are useful as components of grotein CC chips. The nucleic acid sequences of the invention can be used in gene CC therapy. This polynucleotide sequence represents a tumour suppression CC related human fukutin oligonucleotide of the invention.
   Best
                                    Query Match
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                                                                                                      Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 233; 720pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and transfected cells.
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       Local
   Similarity
                                                                                                      BP; 6 A; 3 C; 4 G; 4 T; 0 U; 0 Other;
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Pred.
                                        Score 14.4; DB 1;
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   6.2e+02
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RESULT 990
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CC related human fukutin oligonucleotide of the invention
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                                                                                                                                                        Query Match
Best Local Similarity
Matches 15; Conser
 ABZ64551 standard; RNA; 17
                                                                                                                                                                                                                                    Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 172; 720pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
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                                                                                                         ATTITATTTTCCCTT 1363
                                                                                          ATCTTTATTTTCCCTT 17
                                                                                                                                                                                                                                      BP; 2
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                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                    A; 4 C; 1 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                             1.0%;
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                                                                                                                                                              0;
                                                                                                                                                                             Score 14.4;
Pred. No. 6.
                                                                                                                                                                Mismatches
                                                                                                                                                                             6.2e+02
                                                                                                                                                                                                    DB 1;
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                                                                                                                                                                                                Length
                                                                                                                                                                Indels
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RESULT 991
ACD60861/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HERZ, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or pancreatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ6216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human riborates.
Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis, antisense; enzymatic nucleic acid; hammerhaed ribozyme; DNazyme; inozyme; zinzyme amberzyme; G-cleaver ribozyme; decoy molecule; aptamer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
                                                                            HCV DNAzyme
                                                                                                                                          ACD60861;
                                                                                                                                                                      ACD60861 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                            Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page 133; 185pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAY-2001; 2001US-0294140P.
06-JUN-2001; 2001US-0296249P.
10-SEP-2001; 2001US-0318471P.
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                                                                                                           24-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                             ribozymes of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-140484/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-MAY-2002;
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                                                                                                                                                                                                                                                                                   469
                                                                                                                                                                                                                                                     17 GGGGCGCGCGGCTGCC 2
                                                                                                                                                                                                                                                                                                                   15;
                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                   GGGGCGCGCCGGCTGAC 484
                                                                            substrate sequence #2047.
                                                                                                                                                                                                                                                                                                                                                                            BP; 1 A; 9 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002WO-US016840
                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                           invention
                                                                                                                                                                                                                                                                                                                               1.0%;
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                                                                                                                                                                                                                                                                                                                               Score 14.4; DB 1;
Pred. No. 6.2e+02;
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                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                             Length
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RESULT 992
ADB44694
ID ADB44694
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                                                                                                                               Query Match
Best Local S
Matches 15
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08-JUN-2001;
08-JUN-2001;
24-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
                                                                                                                                                                                      Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 270; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel compound useful for treating cirrhosis, liver failure, hepatocellular carcinoma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blatt
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(BLAT/)
(MACE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-2002; 2002WO-US009187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Draper
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(PAVC/)
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                                                                                                    961
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ROBERTS E.
                                                                                                                                            Similarity
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MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RIBOZYME
BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PAVCO P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LEE P.
                                                                                                    TCGCGCGGCCCCGGCG 976
                                                                       TCGCGCGCCACCGGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Macejak D,
Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; 2001US-00817879.
; 2001US-00877478.
; 2001US-0296876P.
; 2001US-0335059P.
                                                                                                                                                                                      BP; 1 A;
                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PHARM INC
                                                                                                                                            1.0%;
                                                                                                                                                                                      7 C; 7 G; 0 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mcswiggen
                                                                         N
                                                                                                                               0;
                                                                                                                                           Score 14.4; DB 1.
Pred. No. 6.2e+02
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                                                                                                                                Mismatches
                                                                                                                                                                                      2 U; 0 Other;
                                                                                                                                                          DB 1;
                                                                                                                                                          Length 17
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standard;

DNA; 17

27-JUN-2003

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RESULT 993
ACC54422
ID ACC544
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AC ACC544
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AC ACC544
XX
DT 27-JUN
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Consequences, of nucleotides and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as containing the vectors), the encoded polypeptides and antibodies containing the vectors), the encoded polypeptides and antibodies of viral infections or diseases characterized by development of tumours containing the vectors containing the vectors containing the vectors containing the vectors of seases characterized by development of tumours containing the vectors of the nucleotides can be used for diseases and/or prognosis of these diseases. The nucleotides and polypeptides can be used to screen for their specific interactions molecules, containing the nucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss;
primer; probe; tumour suppression; tumour reversion; apoptosis;
virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17 BP; 2 A; 4 C; 1 G; 10 T; 0 U; 0 Other;
                                                                ACC54422;
                                                                                                                         ACC54422
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                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                      ATTTTATTTCCCTT
                                                                                                                                                                                                                                                                                   ATCITATTTTCCCTT 17
                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%;
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                                                                                                                            17
                                                                                                                                                                                                                                                                                                                                      1363
                                                                                                                         BP
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Pred. No. 6.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 17;
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RESULT 994
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XX AD1848
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XX S8; en
KW HCV IN
XX HEQUIN
XX H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence represents an isolated nucleic acid sequence associated with tumour suppression or regression, apoptosis or virus resistance. Invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. The invention is used to diagnose or treat viral disease or disease characterized by development of tumour cells or cellular degeneration
                                                                                                                                                                                                                                                                                                                                                                                                               HCV DNAzyme substrate sequence #2047.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI84801;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid sequences associated with tumor suppression, regression, apoptosis or virus resistance are useful to diagnose and treat viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tuijnder M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-JUN-2001; 2001FR-00008139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-JUN-2001; 2001FR-00008139
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  (BLAT/)
                                                                                                                                                                                                                                 US2003125270-A1
                                                                                                                                                                                                                                                                                    Hepatitis C virus.
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                                                                             18-DEC-2000;
                                                                                                                                18-DEC-2000; 2000US-00740332.
                                                                                                                                                                                   03-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumour suppressor; antitumour; cytostatic; tumour suppression; our regression; apoptosis; virus resistance; diagnosis;
                                                                                                                                                                                                                                                                                                                                    enzymatic nucleic acid; RNA cleavage; hepatitis infection; type I interferon; DNAzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTTTTATTTTCCCTT 1363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           development of tumor cells and cell degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATCTTTATTTTCCCTT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.0%;
illarity 93.8%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 2 A; 4 C; 1 G; 10 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Telerman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    suppressor
                                                                             2000US-00740332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ۶
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.4; DB 1;
Pred. No. 6.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    æ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                    virus;
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RESULT 995
AAA92546/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Enzymatic nucleic acid molecules which specifically cleave from hepatitis C virus (HCV), useful for the treatment of H especially in combination with type I interferon therapy.
The present sequence is one of a large number of antisense oligonucleotides which is directed against one of four human receptor RNA activator (SRA) nucleic acid sequences. Two seri
                                                                         Antisense compounds targeted to steroid receptor RNA activator useful diagnosis, prophylaxis and treatment of diseases associated with the steroid activator, such as infection, inflammation or tumor formation.
                                                                                                                                                                                   (ISIS-)
(BAYU )
                                                                                                                                                                                                                         29-MAR-1999;
                                                                                                                                                                                                                                                                                                       US6107092-A
                                                                                                                                                                                                                                                                                                                                                                        Human;
                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide ISIS# 30213
                                                                                                                                                                                                                                                                                                                                                                                                                          04-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA92546;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 2047; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-031273/03
                                                                                                                                WPI; 2000-586211/55.
                                                                                                                                                          Cowsert
                                                                                                                                                                                                                                                   29-MAR-1999;
                                                                                                                                                                                                                                                                             22-AUG-2000
                                                                                                                                                                                                                                                                                                                                                          SRA inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA92546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ROBE/) ROBERTS E. (PAVC/) PAVCO P A. (MACE/) MACEJACK D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             961 TCGCGCGGCCCCCGGCG 976
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17
                                                                                                                                                                                                                                                                                                                                                          SRA; steroid receptor RNA activator; cytostatic; antiinflammatory; hibitor; cancer; infection; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 l Similarity
15; Conserv
                                                                                                                                                          <u>;</u>
                                                                                                                                                                                   ISIS PHARM INC.
BAYLOR COLLEGE MEDICINE.
                                                    Co1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCGCGCGGCACCGGCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 1 A; 7 C; 7 G; 0 T; 2 U;
                                                  41; 47pp;
                                                                                                                                                          Bennett
                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry
                                                                                                                                                                                                                           99US-00280409
                                                                                                                                                                                                                                                   9905-00280409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.0%;
                                                                                                                                                          Ę,
                                                    English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Roberts
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                                                                                                                                                          O'malley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14.4; DB 1;
Pred. No. 6.2e+02;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'n
                                                                                                                                                          BW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pavco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Macejack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
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uman steroid
series of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e RNA derived
HCV infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         infections,
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RESULT 996
AAH44097
ID AAH440
XX AAH440
XX Oryza
KW Modifi
KW PCR pr
XX Oryza
KW MOZ001
XX ORYZA
XX ORYZA
XX ORYZA
XX ORYZA
XX ORYZA
XX WOZ001
XX OROZO
PD 14-JUN
XX OROZO
PD 14-JUN
XX OROZO
PD 14-JUN
XX OROZO
PR 10-DEC
XX (NORQ
XX ONAShi
PI Hiraga
XX WPI; 2
YX OROZO
PT Charac
XX WPI; 2
YX Tice u
PT Charac
YX WPI; 2
XX OROZO
PT Charac
YX Exampl
XX Exampl

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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                          The present invention describes a set of peroxidase genes found in plants, especially rice, and their homologues, modified forms and fragments, where the second the peroxidase genes in the set are given in AAH44071 to AAH44091. Also described are: (1) promoters for the control of the gene set; (2) the preparation of cassette vectors using the genes and promoters; (3) analysis of plant characteristics using the peroxidase set by isolating RNA from the plant, binding the RNA to a membrane, mixing with a labelled peroxidase gene set, incubating, and detecting the label signal to show which genes in the set are expressed in the sample plant; and (4) DNA microarrays for peroxidase gene expression analysis. The set of genes are used for the analysis of the pattern of peroxidase gene expression in particular rice plants and their component tissues and under different environmental conditions, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ohashi Y,
Hiraga S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -30 oligodeoxynucleotides with a phosphorothioate backbone. The second series comprised chimeric oligonucleotides composed of a central gap region, consisting of ten 2'-deoxynucleotides which was flanked on both sides by four-nucleotide wings. The wings were composed of 2'-methoxyethyl (2'-MOE) nucleotides. Both series contained the same nucleotide sequences. The antisense compounds are useful for research, inflammation or tumour formation. Therapeutically the oligonucleotides are highly safe and are effectively administered to humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oryza sativa; rice; peroxidase; POX; characteristic; gene expression; modification; plant; bacterial infection; Magnaporthe grisea; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Page 62; 258pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New set of rice peroxidase rice under varying condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-DEC-2000; 2000WO-JP008728.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oryza sativa peroxidase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH44097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH44097 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18 BP; 4 A; 4 C; 1 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense oligonucleotides were synthesised. The first series
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NORQ ) JAPAN MIN AGRIC FORESTRY & FISHERIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1491
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACATTAATTGCAGAAA 1506
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          conditions and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer/probe SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sasaki T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genes
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Pred. No. 5.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for analysis of peroxidase expression in production of rice plants with desired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nagamura
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NO:48
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rice plants to provide desired specificities

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ABS52682/
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CC system comprising translation factors and transfer ribonucleic acidd (CRNA) species which translate exogenously added messenger RNA (mRNA) CC with highly selective incorporation at each codon to form a peptide or a peptidominetic product when the system includes one or more tRNA species CC charged with a synthetic amino acid or amino acid analogue. The CC peptide or protein ligands or catalysts, such as insulin, growth hormone CC peptide or protein ligands or catalysts, such as insulin, growth hormone CC peptide or protein ligands or catalysts, such as insulin, growth hormone CC peptide system that offers potentially improved routes to all CC peptides and proteins currently synthesised by alternative routes. This covercomes the limitations of the prior art, e.g. difficulty in CC maintaining purified components and trace contaminants or inefficient CC processivity. There are several advantages associated with performing post-translational modification of peptides, lack of proteases which CC often cause protein display in a pure system, such as an expected lack of contaminants in the selection steps. The present sequence represents a contaminants in the selection steps. The present sequence represents a contaminants of the protein steps. The present sequence represents a contaminants in the selection steps. The present sequence represents a contaminants in the selection steps.
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A new reconstituted cell-free translation system comprising translation factors and tRNA species capable of translating exogenously added mRNAs, useful for the synthesis of peptides or protein ligands or catalysts,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peroxidase gene expression to impart particular characteristics to the plants such as response to bacterial infection by Magnaporthe grisea. present sequence represents a PCR primer/probe for a rice peroxidase (POX) gene, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Translation; ss; splint; cell-free translation system; growth hormone; erythropoietin; ribosome display; mRNA
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                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel reconstituted cell-free translation
                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 15; 83pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Forster AC,
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Pred. No. 5.9e+02
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RESULT ADDITION OF THE RESULT AND ADDITION OF THE RESULT ADDITION OF
                                                CC ontacting a target nucleic acid having one or more sequences selected from 5 3581 base pair sequences in a biological sample with at least one creagent or a series of reagents. The method is useful for detecting the contacting the subject. The set of the 5' and promoter region of the gene DD3 within a contact of the set of oligonucleotides comprising at least three of the coligoners is useful for detecting the complementary sequences. The set of oligonucleotides comprising at least three of the complementary sequences. The set of oligoners is also useful for detecting the set of oligonucleotides within SEQ. ID NO. 1-5 and its complementary sequences. The set of oligonucleotides within SEQ ID NO. 1 and its complementary sequences. The set of all cpd dinucleotides within SEQ ID CC amplification of DNA sequences selected from SEQ ID NO. 1-5 and its complementary sequences selected from SEQ ID NO. 1-5 and its comprising at least one nucleic acid. The methods, nucleic acids, coligonucleotide or PNA-oligomer, kit, array or the set of oligonucleotides is useful for the characterization, classification, classification, sequences, or the predisposition to cell proliferative disorders. The present sequence represents a detection oligonucleotide of the therapy of cell proliferative disorders. The present sequence represents a detection oligonucleotide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting methylation of 5' and promoter region of DD3 gene for diagnosing proliferative disorders comprising contacting target acid with a reagent that distinguishes between methylated and no methylated CpG dinucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to detecting the methylation state of the
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ilarity 93.8%;
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                         CC This invention relates to the identification of a novel susceptibility CC locus AST-1 for asthma and other IgE mediated diseases mapped to the CC human chromosome 7p14-p15. Specifically, it refers to two overlapping CC genes namely GPRA (G-protein coupled receptor for asthma susceptibility) and AAA1 (asthma associated alternatively spliced gene 1). The present CC invention describes identifying single nucleotide polymorphisms, as well as insertion or deletion polymorphisms, occurring at different positions CC in the AST-1 locus, and furthermore providing vectors, host cells, CC primers and probes in order to determine the status of an individual. CC asthma, chronic obstructive pulmonary disease or cancer and other IgE mediated diseases including rhinitis and dermatitis, such that derived CC pharmaceutical compositions exhibit cytostatic and antiasthmatic CC activities. Furthermore, it provides a transgenic animal comprising the asthma locus-1 (AST-1) DNA. This oligonucleotide sequence is a 5 splice junction of the human AAA1 gene, given in Table 11 of the invention.

∴ Matches

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-DEC-2002; 2002US-0435846P.
03-JAN-2003; 2003US-0437895P.
26-MAR-2003; 2003US-0458767P.
09-JUL-2003; 2003US-0486000P.
                                                                                                                                                                                                                                                                                                                         New GPRA polypeptides, useful in preparing a composition for diagnosing, treating or preventing asthma, other IgE-mediated disease, chronic obstructive pulmonary disease or cancer.
                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-500286/47.
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Matches 15
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                                                                                                                                   New hooked or semi-hooked oligonucleotides (see AAQ68869-71, AAQ68873, AAQ68875, AAQ68877, AAQ68879 and AAQ68880) are useful as therapeutic antisense molecules for treating dermatological disorders (e.g. atopical dermatitis, psoriasis, melanoma, T cell lymphoma etc.) viral infections (e.g. herpes simplex, papilloma, hepatitis or HIV); or cancer (when directed against an oncogene), due to their ability to hybridise with target nucleic acid. They can be used ex vivo, e.g., to treat bone marrow grafts. They can also be used for diagnosis or in cosmetics e.g. to block mRNA coding proteins involved in the ageing process such as collagenase or elastase. This linear antisense oligonucleotide is used as a control to see whether the hooked and semi-hooked oligonucleotide exhibit a greater resistance to exonucleases than linear oligonucleotides. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide; antisense; self paired; nuclease resistant; dermatological disorders; viral infection; cancer; atopical derm psoriasis; melanoma; T cell lymphoma; herpes simplex; papilloma; hepatitis; HIV; human immunodeficiency virus; oncogene; collagen.
                                                                                                                                                                                                                                                                                                                                                                                      New oligo:nucleotide(s) self paired at one or both ends - have improved resistance to nuclease(s) and reduced toxicity, useful as anti:sense molecules for treating dermatological disorders, virus infections,
                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                Example 1; Fig 4a; 40pp; French.
                                                                                                                                                                                                                                                                                                                                                                            cancer, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vasseur M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FR2703053-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   elastase; bone marrow graft; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
31-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ68872;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ68872 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEST ) GENSET.
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                                                     l Similarity
15; Conserv
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                    TCTATTTTTTTATTT 1317
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                                                                                                            ₿P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Blumenfeld M,
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                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SA12/m1) used as
                                                                                                            A; 1 C; 2
                                                                  1.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Meguenni
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ΒP
                                                     <u>,</u>
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                                                                   Score 14.4;
Pred. No. 5.
                                                                                                            G; 1 T; 0 U; 0 Other;
                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     control in antisense therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ູດ
                                                                5.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Poddevin
                                                                                DB 1;
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                                                                             Length 20;
                                                        Indels
                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dermatitis;
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                                                      Gaps
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RESULT 1001
AZC03168/c
ID AZC0316
XX AZC0316
XX AZC0316
XX Vaccine
XX Synthet
OS Synthet
OS Chlamyd
XX Synthet
OS Chlamyd
XX CLLamyd
AAX97150
ID AAX9
XX AAX9
AC AAX9
XX I3-9
DT 13-9
XX PCR
XX PCR
XX Resg
                                                                                                                                                                                                                                                                                           RESULT 1002
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17-DEC-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY36754-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, penidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis, pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vaccine; eye disease; conventional trachoma; nonendemic trachoma; paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
Chlamydia trachomatis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ03168 standard;
                                                        PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                             AAX97150;
                                                                                                                                                                                                                                  AAX97150 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 7 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 1584; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genome sequence of Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-371125/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ03168;
                                                                                                                    13-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9928475-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                            1182 TGACGCCGATTCGTGT 1197
                                                                                                                                                                                                                                                                                                                                                                                  19 TGATGCCGATTCGTGT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                 (first entry)
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97FR-00016034.
98US-0107077P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                  ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.4; D
Pred. No. 5.3e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis;

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AAA55806
AAAA55806
XX
XX
AC AAAA1
AC AAA1
AC AAAA1
AC AAA
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Best Local
                                                                                                                                                                                                                                                                                                              Human, DNA methyltransferase; DNA MeTase; antisense oligonucleotide; modulation; inhibition; gene expression; combination therapy; p16; histone deacetylase; HDAC; thymidylate synthase; tumour suppressor; methylation; gene therapy; tumour; cytostatic; antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAY34584-AAY35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human histone deacetylase HD2 antisense oligonucleotide SEQ ID NO:51:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA55806
     19-OCT-1998;
                                                                                                                                                                                                                                                                                         antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA55806 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX91991-X97517 represent PCR primers used to amplify open reading frame and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX9199). C. pneumoniae causes respiratory disease such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genome sequence of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-357842/30
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04-NOV-1998;
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                                                           19-OCT-1999;
                                                                                                                                                                           WO200023112-A1
                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    epitope of C. pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 1881; Disclosure; 1912pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      725
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     98US-0104804P
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98US-0107078P.
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                                                           99WO-US024278.
                                                                                                                                                                                                                                                                                         inflammation;
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Pred. No. 5.3e+02;
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AX (WETH-) METHYLGENE INC.

XX (PI) 2000-339532/29.

XX Inhibiting gene expression e.g. DNA methyltransferase, by treating cells are the symergiatic amount of antisense oligonucleotide and protein of e.g. tumors.

XX Inhibiting gene expression e.g. DNA methyltransferase, by treating cells are the symergiatic amount of antisense oligonucleotide which inhibits the symergistic amount of an artisense oligonucleotide which inhibits capacity and space are product of the gene. Also described are: (1) a method for the gene, and an effective synergistic amount of a protein effector of a product of the gene. Also described are: (1) a method for comprising an antisense oligonucleotide which inhibits expression of the gene in operable association with a protein effector of a gene in a method for comprising an antisense oligonucleotide which inhibitor of a gene comprising an antisense oligonucleotide which inhibitor of a gene for an antisense oligonucleotide which inhibitor of a gene product; (2) a method for inhibiting tumour growth in mammal; (2) an inhibitor of a gene product; (3) and (4) a pharmaceutical composition comprising the inhibitor of the comprising an antisense oligonucleotide which inhibitor of a gene product; (3) and (4) a pharmaceutical composition comprising the inhibitor of (3). The comprising and compositions and comprision to far gene product; (4) and the tinhibit comprises and compositions of the invention that inhibit compation or activity of a gene or gene product may be used to treat the partients having, or predisposed to developing, a disease responsive to comprision of the gene. These may also be used to activate silenced genes to provide missing gene functions and improve a given condition.

CC mathods inhibitors and compositions are useful as probes of the predictivity of a gene product in an experimental cell culture of the product of the predictivity of the predictivity of the predi
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AAZ7563
ID AAZ7563
XX AAZ7563
AC AAZ7563
XX IO-SEP-
XX Human b
XX Human p
XW Human genomic
XW haploty
XW haploty
XW haploty
XW haploty
XW diagnos
XX Homo sa
XX Homo sa
XX WO99545
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PD 28-OCT-
PF 21-APR-
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                                                                                                                                      Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP; PCR primer;
              21-APR-1999;
                                          28-OCT-1999.
                                                                      WO9954500-A2
                                                                                                                           diagnosis; ss.
                                                                                                                                                                                                         Human biallelic marker downstream amplification primer
                                                                                                                                                                                                                                      10-SEP-2001
                                                                                                                                                                                                                                                                                           AAZ75633 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                  ocal
                                                                                               sapiens
                                                                                                                                                                                                                                                                                                                                                                                          726
                                                                                                                                                                                                                                                                                                                                                                                                                     15;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                 99WO-IB000822.
                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Pred. No. 5.3e+02
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RESULT 1005
ABL57552
ID ABL5755
XX ABL5755
XX ABL5755
XX Synthet
XX Concent
KW Concent
KW Polymex
XX Synthet
XX Synthet
XX Concent
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XX Synthet
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ65654 to AAZ69578 represent human biallelic markers from the present Invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ7440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from bharmaceutical agents acting on a disease as well as other treatment.

N.B. The SEQ ID NOS 2822, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not accually given a sequence in the Sequence Listing from the
                               WPI; 2000-657765/64
                                                                                                                                                                                                                                                            20-APR-1999;
                                                                                                                                                                                                                                                                                                              20-APR-2000; 2000EP-00108643
                                                                                                                                                                                                                                                                                                                                                                 25-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                   EP1046717-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Concentration; quantification; mutation detection; polymorphic; polymerase chain reaction; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic deoxyribooligonucleotide poly s.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL57552;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABL57552 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-013267/01.
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23-NOV-1998;
                                                                                                                                                       (NIBI-) JAPAN BIOINDUSTRY ASSOC. (AGEN ) AGENCY OF IND SCI & TECH (KANK-) KANKYO ENG CO LTD.
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nes 15; Conserv
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                                                                               Kanagawa T,
Furusho K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                              99JP-00111601.
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98US-0109732P.
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LTD.
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                                                                                                       Kamagata Y,
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Pred. No. 5.
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                                                                                                          Yamada K,
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                                                                                                          Yokomaku
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Determining the concentration of a target nucleic acid, useful e.g. for detecting genetic mutations, comprises using a fluorescently labeled probe in which emission is reduced by binding to the target nucleic acid.
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Example 6; Page 23; 55pp; English.

CC The invention relates to the determination of the concentration of a CC nucleic acid target, using a fluorescently labeled probe which produces CC reduced fluorescence emission when hybridised to the target nucleic acid. The method comprises measuring the reduction in emission caused by CC hybridisation. The new method is particularly used to quantify target nucleic acids by a real-time polymerase chain reaction, e.g. for CC quantifying microbial cells in co-cultures or symbiotic systems, for CC quantifying microbial cells in co-cultures or symbiotic systems, for CC curves of target nucleic acids to determine a Tm value. Methods of the CC curves of target nucleic acids to be quantified quickly, easily and accurately. Particularly there is no need to remove unbound probe, and no CC materials are introduced that inhibit amplification by Taq polymerase (so CC conventional PCR conditions can be used). The specificity of PCR is kept CC injeh (amplification of primer dimers is delayed), and the limit of CC quantitation is reduced. Complex probes are not needed, and amplification CC can be monitored in real time. The working graph for data analysis CC (automatically generated by a computer) has a higher correlation is CC coefficient than conventional graphs so more accurate quantitation is CC deoxyribooligonucleotide that was used for investigating the effects of the number of G(S) in each target nucleic acid probe its corresponding invention nucleic acid probe

Sequence 20 BP; 4 A; 2 C; 0 G; 14 T; 0 U; 0 Other;

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8
                           Query Match
Best Local
                   Matches
1345 TATATTTTTATTTTCC 1360
                   l Similarity
                   Conservative
                          1.0%;
                   0
                           Score 14.4; DB 1;
Pred. No. 5.3e+02;
                 Mismatches
                                   DB 1;
                                  Length 20;
                    Indels
                    0;
                    Gaps
                    0
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s

TATATTTTTTTTTCC 20

AAH43116 standard; DNA; 20 ВP

AAH43116;

19-SEP-2001 (first entry

Antisense oligo, target HDAC-2 121-141.

cell proliferation; ca fungal infections; ss. Antisense; cell prolif histone deacetylase; HDAC-1; feration; cancer; restenosis; HDAC-2; HDAC-4; inhibitor; psoriasis; protozoal infec infection,

Synthetic

WO200138322-A1.

31-MAY-2001.

22-NOV-2000; 2000WO-IB001881.

23-NOV-1999; 99US-0167035P

(METH-) METHYLGENE INC.

Ö Ruel R, Lavoie R, Thibault C, Abou-Khalil

H

WPI; 2001-432601/46.

ARAH43116
AAH43111
XX
AAH4311
AC
AAH4311
XX
ARAH4311
XX
ARAH411
XX
ARAH41
XX
ARAH4311
XX
ARAH411
XX
ARAH41
X New inhibitors of histone deacetylase e.g. N-hydroxy-5-(4-(benzenesulfonylamino)-phenyl)-4-yn-2-pentanamide for trea for treating cancer,

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RESULT 1007
AAC89545
ID AAC8954
XX AAC8954
XX AAC8954
XX DH HUMAN H
XX Histone
KW Histone
KW HOAC-D;
KW Gene th
XX WO20007
XX WATH-|
XX MATH-|
XX MACleot
PT Antise
PT Macleot
TX WPI, 20
XX
TX WPI, 20
XX
TAB
CC The Dr
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Best Local Similarity
Thes 15; Conserv
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Matches 14; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The sequences given in AAH43115-21 are oligonucleotides which are antisense to the histone deacetylase gene, HDAC-2. These oligonucleotides may be used in combination with an inhibitor of histone deacetylase enzyme function, to given an improved inhibitory effect, thereby reducing the amount of inhibitor required to obtain a given inhibitory effect. Compounds containing these oligonucleotides may be used to treat cell proliferation conditions such as cancer, restencess or psoriasis. They can also be used to treat protozoal and fungal infections
                                                                                                                                       The present invention provides inhibitors of histone deacetylase enzymes such as HDAC-1, HDAC-2, HDAC-3, HDAC-4, HDAC-5, HDAC-C and HDAC-D. These inhibitors may be antisense strands or they may be compounds identified by contacting the enzyme with the compound and measuring the resulting enzyme activity. These inhibitors are useful for treating cancers and for
                                                                                                                                                                                                                                                                  Example 1; Page 24; 125pp; English.
                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide that inhibits expression of a histone deacetylase, useful for treating and/or alleviating the symptoms of neoplasia, or for inhibiting neoplastic cell growth in an animal.
                                                                                                                                                                                                                                                                                                                                                                                           WPI, 2001-016407/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Macleod AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Histone deacetylase; HDAC-1; HDAC-2; HDAC-3; HDAC-4; HDAC-5; HDAC-C; HDAC-C; HDAC-D; cell cycle; tumourigenesis; cancer; inhibitor; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human HDAC-2 antisense sequence SEQ ID NO: 15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-MAR-2001
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                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (METH-) METHYLGENE INC.
                                                                                                                    dentifying which histone deacetylase is involved in a neoplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            726 TGCTGTTGCTGCTGCC 741
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                                                                                ВÞ;
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  Conservative
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                1.0%;
87.5%;
                                                                                7 C; 7 G; 4 T; 2 U; 0 Other;
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1;
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                   Score 14.4; DB 1;
Pred. No. 5.3e+02;
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Pred. No. 5.3e+02;
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    Mismatches
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Homo sapiens.

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RESULT 1008
RESULT 1008
RAC8953
AC AAC8953
XX AAC8953
XX Histone
XX Histone
XX Histone
XX Histone
XX Homo Ba
XX WO20007
XX Gene th
XX
YOU
XX WO20007
XX WALLeod
XX WPI; 20
XX Disclos
XX Di
AAH91976/c
ID AAH9197
XX AAH9197
XX AAH9197
XX AAH9197
XX D9-OCT-
DT 09-OCT-
XX Human 1
XX Human;
KW Human;
KW Bingle
KW chromos
XX OS Homo sa
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Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antisense oligonucleotide that inhibits expression of a histone deacetylase, useful for treating and/or alleviating the symptoms neoplasia, or for inhibiting neoplastic cell growth in an animal
                                                   Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test; chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene therapy; PCR
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                                                                                                                                                                           Human inflammatory bowel disease associated polymorphic site #1051.
                                                                                                                                                                                                                                        09-OCT-2001
                                                                                                                                                                                                                                                                                                  AAH91976;
                                                                                                                                                                                                                                                                                                                                                            AAH91976 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-MAY-1999;
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15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 0 A; 7 C; 7 G; 6 T; 0 U;
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Pred. No. 5
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                                                                                                                      Recognition system; screening; identification; pharmaceutical; toxin; plant protection agent; toxin; venom; carcinogen; venom; teratogen; herbicide; fungicide; pesticide; beta-actin; human; ss.
                             25-MAY-1999;
                                               25-MAY-1999;
                                                                  30-NOV-2000
                                                                                    DE19923966-A1
                                                                                                     Homo sapiens.
                                                                                                                                                            Human beta-actin derived oligonucleotide #6
                                                                                                                                                                              21-MAR-2001
                                                                                                                                                                                                 AAC82913;
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                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 10 A; 3 C; 2 G; 4 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Testing for the bowel disease,
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10-APR-2000; 2000US-0196046P.
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                                                                                                                                                                                                                                                                          TTTATTTTCAGAGACAG 1327
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                                                                                                                                                                                                                                                                                                    ilarity 88.2%;
Conservative
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                              99DE-01023966
                                                99DE-01023966
           RES & TECHNOLOGIES GMBH & CO
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Pred. No. 5.3e+02
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     reverse phase triplex forming oligonucleotide; RP-TFO; protected nucleic acid sequence; PNAS; single nucleotide polymorphism; SNP; short tandem repeat; cancer; Factor V Leiden SNP; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Recognition system, e.g. for identifying nucleic acids, comprises least one recognition unit comprising a region with a defined struadjacent to a region with a randomized structure.
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Pred. No. 5.3e+02;
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                                                                                           The sequence is a second reverse phase triplex forming oligonucleotide, RP-TFO (3' to the SNP) used to analyse Factor V Leiden SNP using the method of the invention. The invention relates to analysing target nucleic acid sequences comprising restricting isolated DNA, hybridising at least one triplex forming oligonucleotide (TFO), adding a 3' to 5' exonuclease to form a protected nucleic acid sequence (PNAS) tail structure, hybridising the captured structure with a single nucleotide polymorphisms (SNP) identification probe and determining the SNP score. The methods can be used for analysing target nucleic acid sequences, specially genomic DNA sequences, to determine if they contain SNPs or use in population genetics, drug development, forensics, cancer, genetic disease research, genomic analysis, diagnostics and therapeutics in
                                                                                                                                                                                                                                                                                                                                                                                          03-NOV-1999;
21-DEC-1999;
07-JUL-2000;
                                                                                                                                                                                                                                                                     Analyzing target nucleic acid sequences, useful for population genetics, drug development and diagnosing cancer, comprises hybridizing triple forming oligonucleotide and probe to target sequence.
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                                          Sequence
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  15; Conserv
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OSTE C
                                                                                    plants and animals
                                                                record issued on 09-SEP-2004 :
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RESULT 1013
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ACF57337;
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DT 16-OCT-2003 (first entry)
XX
DT Human atlastin exon 10 intron
XX
KW Human; atlastin; chromosome 1
KW HSP; neuroprotective; gene th
XX
OS Homo sapiens.
OS Synthetic.
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24-AUG-1999;
30-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to a method for measuring nucleic acids using a nucleic acid probe labelled with a fluorochrome. The nucleic acid probe decreases the fluorescence of the fluorochrome when hybridised with a target nucleic acid, the decrease in the fluorescence is measured. The method can be used for measuring a target nucleic acid
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(KANK-)
(KEIZ-)
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                                                                              chromosome 14; 14q22.1; hereditary spastic paraplegia;
tive; gene therapy; intronic splice site; gene; ds.
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Pred. No. 5.3e+02;
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Best Local (
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                                                                                                                                                                                                    Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New atlastin gene, useful for preparing a composition for treating Hereditary Spastic Paraplegia (HSP) or for identifying subjects who have, or at risk of developing, HSP.
 24-APR-2001; 2001US-0286137P
                                      23-APR-2002; 2002WO-US013135.
                                                                            31-OCT-2002.
                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                  Human oligonucleotide sequence.
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12-SEP-2002; 2002US-00242008.
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Pred. No. 5.3e+02
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                                                                                                                                                                                                  Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cycotatic; gene thera; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allerg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                     Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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Pred. No. 5.3e+02;
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23-APR-2002; 2002WO-US013135.

WO200285308-A2

24-APR-2001; 2001US-0286137P

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RESULT 1016
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Best Local
                                                                                                                                                                                                                        Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                      Human oligonucleotide
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  24-APR-2001; 2001US-0286137P
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                                            23-APR-2002; 2002WO-US013135.
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Pred. No. 5
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5, or 3, end genomic flanking
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CC junctions of genes encoding a polypeptide associated with lung and/or cc antinflammatory steroid and ubiquinone. A composition of the invention cc antinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensiave, communosuppressive, and cytostatic activity. The composition may have a cc use in antisense gene therapy. The composition is useful for treating or cytostenting a respiratory, lung or malignant disease or condition, also cor enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of cf. or reducing sensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of adenosine composition inflammation, lung allergies, or a respiratory disease or condition. Clung surfactant in a subject stissue, or treating bronchoconstriction, increasing inflammation, lung allergies, or a respiratory disease or condition. Composition, but was obtained in electronic format directly from WIPO care for which inthe printed constriction, but was obtained on electronic format directly from WIPO
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Sequence
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     BP;
  16 A; 1 C; 1 G; 2 T; 0 U; 0 Other;
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L, Shahabuddin
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Pred. No. 5.3e+02;
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DB 1;

Length

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Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiathmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Human CCR3 oligonucleotide sequence.

24-APR-2001; 2001US-0286137P 23-APR-2002; 2002WO-US013135. WO200285308-A2 Homo sapiens.

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ARESULT 1018
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                                                                                                                                                                                                                            Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiinflammatory; antiallergic; antiasthmatic; pyotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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24-APR-2001; 2001US-0286137P.
                                          23-APR-2002; 2002WO-US013135.
                                                                                     31-OCT-2002.
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cc junctions of genes encoding a polypeptide associated with lung and/or casal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention cc has antiinflammatory, antiallergic, antiathmatic, hypotensive, cc immunosuppressive, and cytostatic activity. The composition may have a cc use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also cc for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels cf, or reducing sensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of adenosine colong surfactant in a subject's tissue, or treating bronchoconstriction, clung inflammation, lung allergies, or a respiratory disease or condition. The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at figuron-int/pub/published_pct_sequences
  Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or corresponding controls.
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Sequence 20 BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; attendid; ubiquinone; antiinflammatory; antialtergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        inflammation;
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Pred. No. 5
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24-APR-2001; 2001US-0286137P 23-APR-2002; 2002WO-US013135 31-OCT-2002 WO200285308-A2 문 S

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RESULT 1020
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antianflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, laminosuppressive, and cytostatic activity. The composition may have a serious respiratory, lung or malignant disease or condition, also or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of of, or reducing sensitivity to adenosine, reducing or depleting levels of or receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung surfactant in a subject's tissue, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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Matches 15
                                                                                                                                                                                                    Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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   24-APR-2001; 2001US-0286137P
                                                                                                                                                                                                                                                                                                                       Human oligonucleotide sequence.
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                                      23-APR-2002; 2002WO-US013135.
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                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel pharmaceutical composition,
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                                                                                                                                                                                         inflammation;
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Sequence

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G; 11 T; 0 U; 0 Other;

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The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of gunctions of genes encoding a polypeptide associated with lung and/or CC massal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC for enhancing the prophylactic or therapautic respiratory effect of an CC antiinflammatory steroid in a subject, for reducing or depleting levels of continuing sensitivity to adenosine, reducing or depleting levels of of receptor, producing bronchodilation, increasing levels of adenosine continuinflammation, lung allergies, or a respiratory disease or condition.

CC lung inflammation, lung allergies, or a respiratory disease or condition. Once: The sequence data for this patent is not represented in the printed confectivation, but was obtained in electronic format directly from WIPO care for wince in family levels of the printed on the printed confective in the pri
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                                                            ftp.wipo.int/pub/published_pct_sequences
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                                   Similarity
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TGGTTAATCTTTTTTT
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Pred. No. 5.
                             Mismatches
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Human; antisense; lung dysfunction; nasal airway dysfunction; antinflammatory steroid; ubiquinone; antinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine seceptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Human oligonucleotide sequence.

HOMO sapiens

WO200285308-A2

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

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                                                                                                                                                                                                                                                                                                                                                                                     Human oligonucleotide sequence
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L, Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                 beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
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Miller S
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surfactant deplet
                                                                   Human; antisense;
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RESULT 1025 ABD29766

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                                                                                                                           CC comprising oligonucleotides, effective for alleviating
CC pronchoconstriction, respiratory tract inflammation, allergies and
CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC oligonucleotides are derived from a gene encoding or regulating
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC device, in separate containers, (b) the oligonucleotides, (c)
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC beta-adrenergic agonist. The composition is useful for preventing or
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC reduce the amount of target polypeptide present in the lungs. The
CC ulmonary obstruction, and/or bronchoconstriction and/or lung
CC with a disease or condition such as pulmonary vasconstriction,
CC inflammation, allergies, asthma, impeded respiration, respiratory
CC transplantation rejection, pulmonary unmonary vasconstriction, pronchitis or cancer.
CC the oligonucleotides into products that free adenosine into the system
CC the oligonucleotides into products that free adenosine into the system
CC considered adenosine content of the anti-sense oligos corresponding to
CC prevent any unwanted effects due to it
                                                                         Query Match
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                                                                                                                         Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15; SEQ ID NO 4055;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bronchodilating agent.
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CC expression of a target polypeptide associated with lung airway or lung content of a target polypeptide associated with lung airway or lung content of a target polypeptide associated with lung airway or lung content of the invention also describes a kit, that comprises: (a) a delivery content of the invention also describes a kit, that comprises: (b) a delivery content of the invention are adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a constant of the composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The gulmonary obstruction, and/or burnfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, confilmmantion, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, confilmmantion, allergies, asthma, impeded respiration, respiratory confilmmantion, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary cransplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content of fifted the content of the sense environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allex reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
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L, Shahabuddin
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(A) or (A) receptors,
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       This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, composition adenosine sensitivity, levels of adenosine (A) or (A) receptors, composition or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating composition or cancer and can be anti-sense to the corresponding mRNA. The composition or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery composition for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, composition has antiallergic, antiinflammatory, antiasthmatic, composition is useful for preventing or creating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to pulmonary obstruction, and/or bronchoconstriction and/or lung. The
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                                                                                                                                                                                                                                                                                                                                                                                                                 Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                                                                                                                   bronchodilating agent.
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ABC ABD30738;

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ABD30738;

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AC ABD30738;

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Human CCR3-derived oligonu
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Human; antisense; bronchoc
KW respiratory tract inflamma
KW analgesic; hypotensive; in
KW beta-adrenergic agonist; r
KW respiratory distress syndi
KW pulmonary transplantation
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AC ABD30738;
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AC ABC-2002; 2002WO-US013:
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23-APR-2001; 2001US-02860
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CBPIG-) EPIGENESIS PHARM
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C1aim 15; SEQ ID NO 1294:
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C1 comprising oligonucleotic
CC comprising oligonucleotic
CC compression of a target per
CC cliponucleotides are der:
CC cypression of a target per
CC dysfunction or cancer and
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     This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptor surfactant depletion or hyposecretion, when administered to a mammal. To oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.
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                                                                                                                                                                                                                                                                  Claim 15; SEQ ID NO 12949; 763pp; English.
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15; Conserv
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, Shahabuddin
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S;
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RESULT 1028
ABD25109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                     Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antisathmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                             23-APR-2002; 2002WO-US013143
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                                                                                                                                                                                           24-APR-2001; 2001US-0286036P
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                                                                                                                                                                                                                                                                                                                                         Homo
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                                                                                                  Li Y,
Tang
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                                                                                                        Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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Pred. No. 5.3e+02;
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Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targual nucleic acids associated with lung airway or lung dysfunction

targeted to

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CC comprising oligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, CC surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, CC analgesic, hypotensive, immunosuppressive and cycostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or readting a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, pulmonary conditions, emphysema, chronic obstructive pulmonary disease, pulmonary conditions present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system constraint and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1029
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                    23-APR-2002; 2002WO-US013143
                                                                                                                                                                                pulmonary transplantation
                                                                                                                                                                                                  emphysema; chronic obstructive pulmonary
                                                                                                                                                                                                                                                                                                                                                            Human stanniocalcin-derived oligo SEQ ID 1314.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABD22302 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1516 AATTAAAAAAAAAAA 1531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                              rejection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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Pred. No. 5.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  due to it
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                                                                                                                                                                                                  disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 20;
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RESULT 1030
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CC pronchoconstriction, respiratory tract inflammation, allergies and
CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC oligonucleotides are derived from a gene encoding or regulating
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC device, in separate containers, (b) the oligonucleotides, (c)
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC treating a respiratory, lung or malignant disease. The administered
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC reduce the amount of target polypeptide present in the lungs. The
CC inflammation, allergies and/or bronchoconstriction and/or lung
CC with a disease or condition such as pulmonary vasconstriction
CC inflammation, allergies, asthma, impeded respiration, respiratory
CC thymidines present in the target RNA serves to prevent the breakdown of
CC thymidines present in the target RNA serves opponding its pulmonary
CC thymidines present in the target RNA serves opponding its pulmonary
CC thymidines present in the target RNA serves to prevent the breakdown of
CC the oligonucleotides into products that free adenosine into the system
CC entered adenosine content of the anti-sense oligos corresponding to
CC thymidines present in the target RNA serves to prevent the breakdown of
CC the oligonal clotides into products that free adenosine into the system
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CC the content of the product of the system
CC the 
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Best Local (
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Miller S
               Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                        AA102454-derived oligonucleotide SEQ ID
                                                                                                                                                                     29-JUL-2004
                                                                                                                                                                                                               ABD27748
                                                                                                                                                                                                                                                       ABD27748 standard;
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Tang
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L, Shahabuddin
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5.3e+02;
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Query Match
Best Local Similarity
Tatches 15; Conserve
RESULT 1031
ADJ59564/c
ID ADJ59564 standard; DNA; 20 BP.
                                                                                                                                                                                                                                                                                                                                C expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.

C The invention also describes a kit, that comprises: (a) a delivery conditions for adding a carrier and for use of the kit. The composition of the invention has antiallargic, antiinflammatory, antiasthmatic, and the result of the invention has antiallargic, antiantory, antiasthmatic, can beta-adrenargic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or reduce the amount of target polypeptide present in the lungs. The composition control target polypeptide present in the lungs. The conformation, allergies and/or bronchoconstriction and/or lung conformation, allergies and/or bronchoconstriction and/or lung conformation, allergies and/or bronchoconstriction and/or lung conformation, allergies, asthma, impeded respiration, respiratory hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary conformation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense to prevent the breakdown of the oligonucleotides into products that free adenosine into the system consumers and invariant and effect dise in the carget and environment and thereby, to prevent and environment and effect dise in the carget and environment and thereby, to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recepts surfactant depletion or hyposecretion, when administered to a mammal oligonucleotides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 15; SEQ ID NO 6760; 763pp; English.
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                                                                                                                                      666 GACTCACCTCTGACGC 681
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, Tang L,
                                                                                                              GACTCACCTCTGTCGC 16
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ilarity 93.8%;
Conservative
                                                                                                                                                                                                                                                                                                                       unwanted effects due to
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L, Shahabuddin S;
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                                                                                                                                                                                                                    Score 14.4;
Pred. No. 5.
                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                             5.3e+02;
                                                                                                                                                                                                                                   DB 1;
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                                                                                                                                                                                                                                   Length 20
                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (A) receptors,
                                                                                                                                                                                                0
                                                                                                                                                                                                Gaps
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RESULT 1032 ADK80880/c ID ADK8088 XX AC ADK8088 XX DT 20-MAY-XX

ADK80880;

ADK80880 standard;

DNA;

20

ВP

20-MAY-2004

(first entry)

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                                      S
                                                                                                                                                                                      The present invention relates to an oligonucleotide anti-sense to e.g., clinitiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. comprising dense to the conjugate of the conjugate of
                                                                                         Matches
                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-JUL-2003; 2003WO-US023509.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide associated to CCR3 #65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-JUL-2002; 2002US-0399076P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2004011613-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ59564;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (EPIG-) EPIGENESIS PHARM INC.
                                           1524 AAAAAAAAGTAAAAGG 1539
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  17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; SEQ ID NO 420; 85pp; English.
                                                                                         15;
                                                                                                               Similarity
  AAAAAAAGTACAAGG
                                                                                                                                                                                      20
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lin S,
                                                                                                                                                                                   BP;
                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       y L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                   2 A; 4 C; 2 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sandrasagra A,
H, Cong H;
                                                                                                             1.0%;
93.8%;
  N
                                                                                         0;
                                                                                                               Score 14.4;
Pred. No. 5
                                                                                                                                                                                   12 T; 0 U; 0 Other;
                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Aguilar D,
                                                                                                                  .3e+02;
                                                                                                                                     DB 1;
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                                                                                                                                     Length
                                                                                            Indels
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                                                                                         Gaps
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Chimeric

phosphorothicate oligonucleotide to target Nav1.3 #8214.

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RESULT 1033
AD045054/c
ID AD04505
XX
AC AD04505
XX
DT 15-JUL-
XX
DT Human o
XX
CCRL; c
KW Human;
KW CCRL; c
KW tryptas
KW tryptas
KW lung di
KW asthma;
KW asthma;
KW asthma;
KW acute x
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Best Local S
Matches 15
Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicary; or ataxia with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target different regions of the human Navl.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               wew antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition
                                                                                                                                                                      Human oligonucleotide #420
                                                                                                                                                                                                             15-JUL-2004
                                                                                                                                                                                                                                                  ADO45054;
                                                                                                                                                                                                                                                                                     ADO45054 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 8214; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-AUG-2002; 2002US-0403416P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 93.0
                                                                                                                                                                                                                                                                                                                                                                                                     1252 TTTTGTTTTTTAATCA 1267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2004-203785/19.
                                                                                                                                                                                                                                                                                                                                                                              17 TTTTGATTTTTAATCA 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention relates to an antisense compound
                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; 2 C; 3 G; 4 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.4;
Pred. No. 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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RESULT 1034 ADO55869/c ID ADO5586

ADO55869 standard; DNA;

20

ВP

AAAAAAAGTACAAGG

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                                                                                                                                                                                                                                                                                             The invention relates to oligonucleotides anti-sense to an initiation CC codon, coding region, 5' or 3' intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II) CF receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC concorded target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The CC concording interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, CCR3, Ectaxin-1, CRANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CRANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CRANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CRANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CRANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CRANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CRANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CRANTES, MCP4, CD23, ICAM, VCAM, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CR1, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CR1, CTYPTASE A, CCR1, CCR3, Ectaxin-1, CR1, CTYPTASE A, CTYPT
                                                                                                        Query Match
Best Local
                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel single or multiple target oligonucleotide anti-sense to e.g. Conitiation codon, intron of respiratory disease-relevant gene e.g. CORANTES, MCP4, useful for prophylaxis or treating respiratory disease
                                                                                                                                                                       Sequence 20 BP; 2 A; 4 C; 2 G; 12 T; 0 U; 0 Other;
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23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                           allergic rhinitis, acute respiratory distress syndrome, pulmonary
hypertension, lung inflammation, bronchitis, airway obstruction o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2; SEQ ID NO 420; 174pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-293804/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NYCE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2004
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                                                                                                                                                                                                                                         bronchoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-JUL-2003; 2003US-00627930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lung inflammation; bronchitis; airway obstruction; bronchoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SHAH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AGUI/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (TANG/)
                                        1524 AAAAAAAAGTAAAAGG 1539
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                                                                                   15,
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SANDRASAGRA
                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MILLER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AGUILAR D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TANG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sandrasagra A, Tang
in S, Lu H, Cong H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ξ
                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ×
                                                                                                        1.0%;
                                                                                                                                                                                                                                       This sequence
N
                                                                                 0
                                                                                                        Score 14.4;
Pred. No. 5
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                                                                                                                                                                                                                                         represents an
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                                                                                                          .3e+02
                                                                                                                               DB 1; Length
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                                                                                                                                                                                                                                         oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene e.g. CCR1,
                                                                                   0,
                                                                                      Gaps
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RESULT 1035
AD055807
ID AD05580
XX
AC AD05580
XC AD05580
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC The present invention relates to antisense compounds targeted to a control of control of the control of con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense therapy, human, NIMA-related kinase 6; never in mitosis gene a-related kinase 6; hyperproliferative disorder; cancer; cytostatic; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotides for modulating never in mitosis, gene a (NIMA)-related kinase 6 expression, useful for diagnosing, preventing or treating diseases associated with the kinase, e.g. hyperproliferative
     Antisense therapy; human; NIMA-related never in mitosis gene a-related kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-NOV-2002; 2002US-00295471.
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                                                                               Human NIMA-related kinase 6 DNA, antisense oligonucleotide
                                                                                                                                                                                 AD055807;
                                                                                                                                                                                                                                  ADOSS807 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-NOV-2002; 2002US-00295471.
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                                                                                                                                12-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                                       1036
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                                                                                                                                                                                                                                                                                                                                                                                                       AGTGGGCGGCGGTGGT 1051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 BP; 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 9 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.4; D
Pred. No. 5.3e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
kinase 6;
6; hyperproliferative disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 20;
                                                                                  #30
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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RESULT 1036
ADP20520/c
ID ADP2052
XX
AC ADP2052
XT 26-AUG-
XX
DE Transcr
XX
CYtOsta
KW cytosta
KW AP-2 al
                                                                                                                                                                                                        멼
                                                                                                                                                                                                                              S
                                                                                                                                                                                                                                                                                                                          The present invention relates to antisense compounds targeted to a nucleic acid encoding human never in mitosis gene a related kinase 6 (NIMA-related kinase 6). The antisense compound comprises an antisense coligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression of NIMA-related kinase 6. The antisense coligonucleotide is a chimeric oligonucleotide. The antisense coligonucleotide is a chimeric oligonucleotide. The antisense coligonucleotide does not necessary to be a comprise at least one modified sugar moiety, preferably a 2'-O-methoxyethyl (2'-MOE) sugar moiety. The antisense oligonucleotide further comprises at least one modified nucleobase, preferably a 5-methylcytosine. The antisense oligonucleotides are useful for the treatment of diseases such as hyperproliferative disorders, e.g. cancer. The present sequence represents an antisense oligonucleotide used in the examples of the
                                                                                                                                                                                                                                                                 Best Loc
Matches
                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI;
                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                      present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotides for modulating never in mitosis, gene (NIMA)-related kinase 6 expression, useful for diagnosing, preventing treating diseases associated with the kinase, e.g. hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dobie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-NOV-2002; 2002US-00295471.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Кeу
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; cytostatic; phosphorothioate;
                                            Transcription
                                                                          26-AUG-2004
                                                                                                     ADP20520
                                                                                                                                 ADP20520 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ISIS-)
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                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                     1036
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                                                                                                                                                                                                                                                                 l Similarity 93.8
15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SISI
                                                                                                                                                                                                                                    AGTGGGCGGCGGTGGT 1051
                                                                                                                                                                                                        AGTGGGCTGCGGTGGT 16
                                                                                                                                                                                                                                                                                                                          ₿₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 44; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002US-00295471.
                                                                        (first
                                            factor AP-2 antisense oligonculeotide segid 67.
                                                                                                                                                                                                                                                                                                                         2 A; 4 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "This oligonucleotide has a phosphorothioate backbone and 2'-methyoxyethyl (2'-MOE) wings at the and 3' ends, which are 5 nucleotides in length at ea end. All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
1. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mod_base= OTHER
                                                                        entry)
                                                                                                                                                                                                                                                                             1.0%;
                                                                                                                                                                                                                                                                                                                          9 G;
                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                               Score 14.4; DB 1
Pred. No. 5.3e+02
0; Mismatches
                                                                                                                                                                                                                                                                                                                          5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88
                                                                                                                                                                                                                                                                                                                          0 Other;
                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      each
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cytostatic; AP-2-Inhibitor-Alpha; AP-2 alpha; AP-2 alpha modulator; AP-2 alpha associated disorder; hyperproliferative disorder; human;

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ADP21858/c
ID ADP2185
XX ADP2185
XX ADP2185
XX ADP2185
XX ADP2185
XX 26-AUG-
XX Cytosta
XW cytosta
XW cytosta
XW ornithi
KW hyperpr
XX
OS Homo sa
XX
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                                                                                                                                                                                                                                                                                                                                                                                     ঠ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention describes a compound (1) 8-80 nucleobases in length CC targeted to a nucleic acid molecule encoding AP-2 alpha. The compound CC specifically hybridises with a nucleic acid molecule encoding AP-2 alpha (1) 96. SEQ ID NO: 4), and inhibits the expression of AP-2 alpha. Also CC comprising contacting the expression of AP-2 alpha in cells or tissues CC comprising contacting the cells or tissues with (1); screening for a CC modulator of AP-2 alpha by contacting a preferred target segment of a CC nucleic acid molecule encoding AP-2 alpha with one or more candidate CC modulators of AP-2 alpha, and identifying one or more modulators of AP-2 alpha, and identifying one or more modulators of AP-2 alpha, and identifying one or more modulators of AP-2 alpha expression, which modulate the expression of AP-2 alpha; a CC diagnostic method for identifying a disease state; and a kit or assay CC device comprising (1). The compound is useful for treating an animal CC having a disease or condition associated with AP-2 alpha, particularly a CC hyperpoliferative disorder. The compounds may be used for diagnostics, CC therapeutics prophylaxis and as research reagents; or as tools in CC differential and/or combinatorial analyses to elucidate expression CC calle and tristing anytion or the entire complement of genes expressed within CC calle and tristing anytion or the entire complement of genes expressed within
                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compounds targeted to nucleic acid molecules encoding AP-2 alphabits the expression of AP-2 alpha, useful for treating AP-2 alassociated disease or condition, particularly a hyperproliferative
                      Homo sapiens
                                                        hyperproliferative
                                                                              cytostatic; gene therapy; ornithine decarboxylase 1; ornithine decarboxylase 1 associated disorder;
                                                                                                                                       Human ornithine decarboxylase 1 primer seqid 6.
                                                                                                                                                                               26-AUG-2004
                                                                                                                                                                                                                   ADP21858
                                                                                                                                                                                                                                                          ADP21858 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 67; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-440306/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-DEC-2002; 2002US-00315962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-DEC-2002; 2002US-00315962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transcription factor; antisense oligonucleotide; antisense technology;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AP-2 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cells and tissues. This sequence represents a human transcription factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2004109848-A1
                                                                                                                                                                                                                                                                                                                                                                                             726
                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                            TGCTGTTGCTGCC 741
                                                                                                                                                                                                                                                                                                                                                      recrectectect 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 5 A; 6 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dean NM,
                                                                                                                                                                             (first entry)
                                                            disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Freier SM,
                                                                                                                                                                                                                                                            ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                            cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.4; DB 1;
Pred. No. 5.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dobie KW
                                                            human; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        alpha and
                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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PT XXX
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                                                                                                                                                                                                                                                                                                                                                     RESULT 1038
                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention describes a new compound, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding ornithine decarboxylase 1, specifically hybridises with the nucleic acid encoding ornithine decarboxylase 1 comprising 2035-bp sequence and inhibits expression of ornithine decarboxylase 1. Also described are: inhibiting the expression of ornithine decarboxylase 1 in cells or tissues; screening for a modulator of ornithine decarboxylase 1; identifying a disease state; a kit or assay device comprising the compound; and treating an animal having a disease or condition associated with ornithine decarboxylase 1. The oligonucleotide compound is useful for preparing a composition for treating hyperproliferative disorder, e.g. cancer. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New oligonucleotide compound that inhibits expression of ornithine decarboxylase 1, useful for preparing a composition for treating hyperproliferative disorder, e.g. cancer.
                        WPI; 1995-018287/03.
                                                                            16-APR-1993;
                                                                                                    16-APR-1993;
                                                                                                                              01-NOV-1994
                                                                                                                                                        JP06303997-A
                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                         aggregate;
                                                                                                                                                                                                                        Analysis;
                                                                                                                                                                                                                                             Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                          04-AUG-1995
                                                                                                                                                                                                                                                                                                  AAQ75552;
                                                                                                                                                                                                                                                                                                                            AAQ75552 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 8 A; 5 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 21; SEQ ID NO 6; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bennett CF, Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-DEC-2002; 2002US-00316244.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-DEC-2002; 2002US-00316244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            decarboxylase 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-)
                                                  (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                     729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2004-440337/41.
                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                                                                                                                                                                                                                                                                                                                15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                              rerrecrecter s
                                                                                                                                                                                                                                                                                                                                                                                                              TGTTGCTGCTGCCTTT
                                                                                                                                                                                                           gene expression; reverse transcription; primer; cDNA;
restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PHARM INC
                                                                                                                                                                                                                                                                           (first
                                                                            93JP-00112515
                                                                                                     93JP-00112515
                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                      744
                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.4;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                            .3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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Analysis of cDNA and gene expression -

by amplification of

mRNA followed

Example

3; SEQ

ij

NO 496; 171pp; English.

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RESULT 1039
ADL79331
XX ADL7933
XX ADL7933
XX ADL7933
XX ADL7933
XX 20-MAY-
XX RNA int
KW RNA int
KW RNA int
KW Short i
KW Short i
KW Short i
KW Short i
KW Cytosta
KW HER2; E
XX HOMO 88
X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           £xgxnnnnnn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                    06-JUN-2002;
06-JUN-2002;
03-JUL-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-FEB-2002;
11-MAR-2002;
29-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene, epidermal growth factor receptor; EGFR; HER2; EGFR2; neu; erbB2; c-erb-B-2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A method for the analysis of cDNA comprises (a) preparing an aggregate (double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) dispesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) slectrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human HER2
                                                           New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADL79331 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                               19-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-FEB-2003; 2003WO-US005045
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 5; llpp; Japanese.
                                                                                                                                                                                                                                                               (RIBO-) RIBOZYME PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1246 TCTTTGTTTTGTTTTTTAA 1264
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (EGFR2) sinA lower strand, SEQ ID NO:496.
                                                                                                                                                                                                                                                                                                                 2002US-0393924P.
2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
2002US-00251117.
2002US-00277494.
2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2002US-00163552.
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2002WO-US016840.
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                                                                                                                                                                                                      Pavco
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                                                                                                                                                                                                                                                               INC.
                                                                                                                                                                                                      Beigelman
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Pred. No. 6.
                                                                                                                                                                                                      ٢
                                                                                                                                                                                                      Fosnaugh K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6.1e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          س
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                                                                                                                                                                                                         Jamison
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CC interference. The siNAs may or may not comprise ribonucleotides and may CC be double or single stranded. They further comprise sense and antisense CC regions, or alternatively are assembled from a sense oligonucleotide and CC an antisense oligonucleotide. Specifically, the siNAs include short CC interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short CC can contain deoxyribonucleotides, and can be unmodified or chemically modified, CC can contain deoxyribonucleotides, and can be chemically synthesised, CC expressed from a vector or enzymatically synthesised. The invention also CC relates to kits for the in vitro or in vivo delivery of siNA; conjugates CC and/or complexes of siNA; and vectors that express siNA. The siNAs are CC used to modulate expression of EGFR genes in cells, tissue explants or CC organisms (e.g., by ex vivo gene therapy), or in grafts and transplants CC organisms (e.g., by ex vivo gene therapy), or in grafts and transplants CC identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide CC identification, and validation, genetic engineering, pharmacogenomics, CC polymorphisms). The present sequence represents the lower strand of a CC (EGFR2)-targeted double-stranded siNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to short interfering nucleic acids (siNA) which downregulate expression of one or more human epidermal growth factor receptor (EGFR) genes (including HER1, HER2 HER3 and HER4) by RNA
  Sequence 19
B₽;
  16
ð
2 C; 0 G; 0 T; 1 U;
  0 Other;
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S Matches Query Match Local 1519 TAAAAAAAAAAAGTAAAA 1537 :||||||| |||| 15; 1 UAAAAAAACAAAACAAAAA 19 Similarity Conservative 1.0%;
78.9%; 1; Mismatches Score 14.2; DB 1; Pred. No. 6.1e+02; ₩ --Length Indels 0, Gaps 0

밁

RESULT 1040
ADL79082/c
ID ADL7908
XX ADL7908
XX ADL7908
XX ZO-MAYXX RNA int
XX RNA int
XX RNA int
XX ENA int 06-JUN-2002; 06-JUN-2002; 03-JUL-2002; 29-AUG-2002; 05-SEP-2002; 09-SEP-2002; 19-SEP-2002; 21-OCT-2002; 20-FEB-2002; 11-MAR-2002; 29-MAY-2002; RNA interference; short interfering nucleic acid; siNA; miRNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; cytostatic; human; oncogene; epidermal growth factor receptor; 20-FEB-2003; 28-AUG-2003. WO2003070912-A2 Human HER2 (EGFR2) transcript target sequence/siNA upper strand, SEQ:247. 20-MAY-2004 ADL79082; ADL79082 standard; RNA; 19 sapiens. EGFR2; 2002US-00163552. 2002US-0386782P. 2002US-0393924P. 2002US-0406784P. 2002US-0406784P. 2002US-04092378P. 2002US-00251117. 2002US-00277494. 2002US-0358580P. 2002US-0363124P. neu; erbB2; c-erb-B-2; target sequence; ss. 2003WO-US005045 2002WO-US016840. (first entry)

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RESULT 1041
ABZ89546/c
ID ABZ8954
XX ABZ8954
XX ABZ8954
XX IT-OCT-
XX Human o
XX Human o
XX Human;
KW Antiast

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                 Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             downregulate expression of one or more human epidermal growth factor receptor (EGFR) genes (including HER1, HER2 HER3 and HER4) by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense
          23-APR-2002; 2002WO-US013135
                                                                          31-OCT-2002
                                                                                                                                           WO200285308-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ89546;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABZ89546 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; SEQ ID NO 247; 171pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-697612/66.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HER2 transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1519 TAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                            inflammation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                         respiratory disease; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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Pred. No. 6.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression of
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e epidermal
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RESULT 1042
ABD25776/c
ID ABD2577
XX ABD2577
XX ABD2577
XX ABD2577
XX ABD2577
XX Human;
XW respirac
XW surfact
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                      Human; antisense; bronchoconstriction; allergy, hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary hyportension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine rung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a novel pharmaceutical composition, which first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 18 A; 0 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       initiation codon, coding region, 5' or 3' end genomic flanking regions' and 3' intron-exon junctions, or regions within 2-10 nucleotides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-APR-2001; 2001US-0286137P
                                                                                                                                                                                               pulmonary transplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AI085559 DNA fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABD25776;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABD25776 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               junctions of genes encoding a polypeptide associated with lung and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (EPIG-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1246 TCTTTGTTTTTTTAA 1264
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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, Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sandrasagra A,
L, Shahabuddin
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es 3;
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WO200285309-A2 Homo sapiens.

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RESULT 1043
AAQ75579
ID AAQ7557
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AC AAQ7557
XC AAQ7557
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DT 04-AUG-
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REVETBO
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REV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC surfactant depletion or hyposecretion, when administered to a mammal. The CC oligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallersic, antiinflammatory, antiasthmatic, and beta-adrenersic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The CC pulmonary obstruction, and/or bronchoconstriction and/or lung confilmmation, allergies and/or bronchoconstriction and/or lung confilmmation, allergies and/or bronchoconstriction and/or lung confilmmation, allergies and/or bronchoconstriction, respiratory distances by an experience of the compositation, respiratory confilmmation, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary confilms present in the target RNA serves to prevent the breakdown of the oligonucleotides into product enterties the protection into the system content of the anti-sense oligos corresponding to the oligonucleotides into product enterties of prevent the breakdown of the oligonucleotides into product enterties of prevent the breakdown of the oligonucleotides into product enterties of the anti-sense oligos corresponding to the oligonucleotides into product enterties of the enterties into the system contents.
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Best Local (
Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                 Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the oligonucleotides into products that e.g., lung, brain, heart, kidney, etc, t prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) rec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 15; SEQ ID NO 4788; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acids associated bronchodilating agent.
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                                                                                                                                                                       04-AUG-1995
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L, Shahabuddin
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Pred. No. 5.8e+02
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Best Local Similarity
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                     Disclosure; Page 5; 11pp; Japanese
                                            Analysis of cDNA and gene expression -
by digestion with restriction enzymes.
                                                                                WPI; 1995-018287/03.
                                                                                                                              16-APR-1993;
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by digestion with restriction enzymes.
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A method for the analysis of cDNA comprises (a) preparing an aggregate

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RESULT 1046
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Best Local Similarity
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                                                                                                                                                                    Sequence
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aggregate; restriction enzyme;
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Pred. No. 5.8e+02
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                                                                                                                                                       The invention relates to a novel pharmaceutical composition, which has a cliret active agent comprising an oligonucleotide antisense to the confirmation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of cylonctions of genes encoding a polypeptide associated with lung and/or constant airway dysfunction and a second active agent comprising an antial ammatory steroid and ubiquinone. A composition of the invention cc has antiinflammatory, antiallergic, antiasthmatic, hypotensive, have a cuse in antisense gene therapy. The composition is useful for treating or compressive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also composition as the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels cf. or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine of lung surfactant in a subject's tissue, or treating bronchoconstriction. CC lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO cat figure.
                                                                                            Query Match
Best Local 9
                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                       Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 3936; 872pp; English.
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L, Shahabuddin
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RESULT 1047 AAQ75596

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                                                                                                                                        aggregate;
                                                                                                                                                                             Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                         04-AUG-1995
                                                                                                                                                                                                                                                           AAQ75597 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 2 A;
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              16-APR-1993;
                                      16-APR-1993;
                                                                                                                                                     Analysis; gene expression; reverse transcription; primer; cDNA;
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Best Local S
Matches 16
                          A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ filles AQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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Sequence 20 BP; 1 A; 1 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                Disclosure; Page 5; llpp; Japanese.
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Query Match

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                                                                                             The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonuclectide antisense to the CC initiation codon, coding region, 5 or 3 end genomic flanking regions, CC 5 and 3 intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of antiinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing sensitivity to adenosine, reducing levels of adenosine or CC lung surfactant in a subject's tissue, or treating bronchoconstriction, CC lung inflammation, lung allegies, or a respiratory disease or condition. CC specification, but was obtained in electronic format directly from WIPO care for wino intronible and not represented in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                           ftp.wipo.int/pub/published_pct_sequences
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L, Shahabuddin
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                       CC composition of igonucleotides, effective for alleviating composition of igonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and composition or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating continged by the control of a target polypeptide associated with lung airway or lung constituction or cancer and can be anti-sense to the corresponding mRNA. CC device, in separate containers, (b) the oligonucleotides, (c) device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallegic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a composition composition composition composition composition composition composition of the administered composition composition composition of the administered composition or availability, or to increase the degradation of the target mRNA or to creduce the amount of target polypeptide present in the lungs. The confidence the amount of target polypeptide present in the lungs. The confidence in and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABD21542 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention
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                                                                                                                                                                                                                                                                                  A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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Pred. No. 5.
                                                                                                                Score 14.2; DB 1;
Pred. No. 5.8e+02;
0; Mismatches 3;
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RESULT 1053

30-SEP-1994

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RESULT 1054
AAQ68872
ID AAQ6887
XX AAQ6887
XX 25-MAR-
DT 31-MAY-
DT 31-MAY-
XX Oligonu
XX Oligonu
KW Oligonu
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Best Local :
                                                                                                       Oligonucleotide; antisense; self paired; nuclease resistant; dermatological disorders; viral infection; cancer; atopical c psoriasis; melanoma; T cell lymphoma; herpes simplex; papillo hepatitis; HIV; human immunodeficiency virus; oncogene; colla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEO files AQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                           elastase; bone marrow graft; ss.
                                                                                                                                                                                   Oligonucleotide (SA12/ml) used as control in antisense therapy
                                                                                                                                                                                                               25-MAR-2003
31-MAY-1995
                                                                                                                                                                                                                                                           AAQ68872;
                                                                                                                                                                                                                                                                                        AAQ68872 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 5; 11pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
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                                 FR2703053-A1
                                                             Synthetic
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(first entry)
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Pred. No. 5.
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RESULT 1055
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New hooked or semi-hooked oligonucleotides (see AAQ68869-71, AAQ68873, AAQ68875, AAQ68877, AAQ68879 and AAQ68880) are useful as therapeutic antisense molecules for treating dermatological disorders (e.g. atopical dermatitis, psoriasis, melanoma, T cell lymphoma etc.) viral infections (e.g. herpes simplex, papilloma, hepatitis or HIV); or cancer (when directed against an oncogene), due to their ability to hybridise with target nucleic acid. They can be used ex vivo, e.g., to treat bone marrow grafts. They can also be used for diagnosis or in cosmetics e.g. to block mRNA coding proteins involved in the ageing process such as collagenase or elastase. This linear antisense oligonucleotide is used as a control to see whether the hooked and semi-hooked oligonucleotides exhibit a greater resistance to exonucleases than linear oligonucleotides. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New oligo:nucleotide(s) self paired at one or both ends - have improved resistance to nuclease(s) and reduced toxicity, useful as anti:sense molecules for treating dermatological disorders, virus infections,
              Fink JK,
                                                                        21-SEP-2001; 2001US-0323997P.
12-SEP-2002; 2002US-00242008.
                                                                                                                    13-SEP-2002; 2002WO-US029165.
                                                                                                                                                                                   WO2003026566-A2
                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                           Human; atlastin; chromosome 14; 14q22.1; hereditary spastic paraplegia; HSP; neuroprotective; gene therapy; intronic splice site; gene; ds.
                                                                                                                                                                                                                                                                                                         Human atlastin exon 10 intronic acceptor splice site.
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                                                                                                                                                    03-APR-2003
                                                                                                                                                                                                                                 Homo sapiens
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                                           (UNMI ) UNIV MICHIGAN
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              Zhao X;
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chromosome 14 (more specifically to 14922.1). Also described: (1) an isolated atlastin polypeptide; (2) identifying subjects who have, or are at risk of developing, hereditary spastic paraplegia (HSP); (3) a kit for determining if a subject has, or at risk of developing, HSP; (4) a computer readable medium encoding a representation of the atlastin nucleic acid sequence or polypeptide; (6) identifying subjects at risk of carrying an allele for HSP, and (7) treating a patient with HSP, Atlastin has neuroprotective activity and can be used in gene therapy. The atlastin nucleic acid is useful for preparing a composition for treating atlastin nucleic acid is useful for preparing a composition for treating HSP or for identifying subjects who have, or at risk of developing, HSP. The present sequence represents an atlastin intronic splice site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New atlastin gene, useful for preparing a composition for treating Hereditary Spastic Paraplegia (HSP) or for identifying subjects who have, or at risk of developing, HSP.
                                                                                                                                                                                                                                                                                                                                                                                                                         Example 6; Page 102; 111pp; English.
                                                                                                                                                                                                                                                                                                                                                                          The present invention describes human atlastin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-371871/35.
                                                                                                                                                                                                                                                                                                                                                                             which is located
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Search completed: November Job time : 22 secs ν, 2004, 12:44:19

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1245 ATCTTTGTTTTTTTTTTA 1263

19

Query Match Best Local Matches

16;

Conservative

0;

Similarity

1.0%;

Score 14.2; DB 1; Pred. No. 5.8e+02; Mismatches

DB 1;

Length 20; Indels

0

Gaps

0

Sequence 20 BP; 2 A; 2 C; 1 G; 15 T; 0 U; 0 Other;

C 1 19.6 1.4 27 1 AZB27502 C 2 18.2 1.3 24 1 AZB27502 C 2 17.8 1.3 25 1 AZB27377 C 17.6 1.3 22 1 AZB21080 C 19 17.6 1.3 22 1 AZB21080 C 10 17.6 1.3 24 1 CZP291636 C 11 17.2 1.2 22 1 AZB20953 C 11 17.2 1.2 22 1 AZB20953 C 11 17.2 1.2 22 1 AZB20958 C 12 17.2 1.2 22 1 AZB20958 C 13 17.2 1.2 22 1 AZB20958 C 14 17 1.2 22 1 AZB20958 C 15 16.8 1.2 20 1 ALD38577 C 2 16 8 1.2 20 1 ALD38576 C 2 16 8 1.2 20 1 ALD38576 C 2 2 16.8 1.2 20 1 ALD38576 C 2 2 16.8 1.2 20 1 ALD38754 C 2 2 16.8 1.2 20 1 ALD38754 C 2 2 16.8 1.2 20 1 ALD38759 C 2 3 16.8 1.2 20 1 ALD38759 C 2 4 16.8 1.2 20 1 ALD38759 C 2 5 16.8 1.2 22 1 ALD38759 C 2 6 16.8 1.2 23 1 CZP39801 C 2 7 16.6 1.2 23 1 CZP39801 C 2 7 16.6 1.2 23 1 CZP39801 C 2 8 16.6 1.2 23 1 CZP39801 C 2 9 16.6 1.2 23 1 AZB4859 C 2 1 16.4 1.2 20 1 AZB4859 C 2 1 16.4 1.2 20 1 AZB4859 C 2 1 16.4 1.2 21 AZB4859 C 2 1 16.5 1.2 23 AZB4859 C 2 1 16.6 1.2 23 AZB4859 C 2 1 16.8 AZB477 C 2 1 16.8 AZB48564 C 2 1 16.8 AZB4777 C 2 1 16.8 AZB48564 C 2 1 16.8 AZB48564 C 2 1 16.8 AZB48564 C	Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. SUMMARIES Result Query No. Score Match Length DB ID Description	<pre>seq length: 8 seq length: 80 seq length: 80 ssing: Minimum Match 0% Maximum Match 100% Listing first 128 summaries rstl2.seq:*</pre>	Scoring table: IDENTITY_NUC Gapop 10.0 , Gapext 0.5 Searched: 126 seqs, 2731 residues Total number of hits satisfying chosen parameters: 252	(without alignments) 3.826 Million cell upda le: US-10-643-432-12 fect score: 1401 unda fect score: 1401	GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd. OM nucleic - nucleic search, using sw model Run on: November 2, 2004, 12:50:31 ; Search time 2 Seconds
C 74 15.6 1.1 22 1 CF33430 CC 74 15.6 1.1 22 1 CF334781 CC 75 15.6 1.1 22 1 CF334781 CC 76 15.6 1.1 22 1 CF336280 CC 77 15.6 1.1 22 1 CF337580 CC 80 15.6 1.1 22 1 CR34550 CC 80 15.6 1.1 22 1 AZ31036 CC 80 15.6 1.1 22 1 AZ31036 CC 80 15.6 1.1 22 1 AZ31636 CC 80 15.6 1.1 22 1 AZ40190 CC 80 15.6 1.1 22 1 AZ40190 CC 80 15.6 1.1 22 1 AZ40190 CC 80 15.6 1.1 22 1 AZ42430 CC 90 15.6 1.1 22 1 AZ42430 CC 90 15.6 1.1 22 1 AZ46360 CC 90 15.6 1.1 22 1 AZ66360 CC 90 15.6 1.1 22 1 AZ6960 CC 90 15.6 1.1 22 1 AZ6960 CC 90 15.6 1.1 22 1 AZ7970 CC 90 15.6 1.1 22 1 AZ89461 CC 100 15.6 1.1 22 1 AZ8946	65 15.6 1.1 22 1 67 15.6 1.1 22 1 67 15.6 1.1 22 1 69 15.6 1.1 22 1 70 15.6 1.1 22 1 71 15.6 1.1 22 1 72 15.6 1.1 22 1 73 15.6 1.1 22 1	56 15.8 1.1 21 1 57 15.8 1.1 22 1 58 15.8 1.1 22 1 59 15.8 1.1 22 1 60 15.8 1.1 22 1 61 15.8 1.1 22 1 62 15.8 1.1 22 1 63 15.6 1.1 22 1 64 15.6 1.1 22 1	15.8 1.1 20 1 15.8 1.1 20 1 15.8 1.1 20 1 15.8 1.1 20 1 15.8 1.1 21 1 15.8 1.1 21 1	16.2 1.2 23 1 16.1 19 1 15.8 1.1 19 1 15.8 1.1 19 1 15.8 1.1 20 1 15.8 1.1 20 1 15.8 1.1 20 1	16.4 1.2 22 1 16.2 1.2 21 1 16.2 1.2 22 1 16.2 1.2 23 1
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ACCESSION
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121
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27 bp DNA linear GSS 20-FEB-20(
2M0103020R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGC2M0103020 R, genomic survey sequence.

AZ827502

AZ827502.1 GI:12997410
                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleoston Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; N. (bases 1 to 27)

1 (bases 1 to 27)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Manoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse, Whole genome scaffolding with paired end reads from 10kb Mouse, Whole genome scaffolding with paired end reads
                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.00
Plate: 0103 row: 0 column: 20
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                 University of Utah Genome University of Utah
                                                                                                                                                                                                                                                                                                           Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
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Location/Qualifiers
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801 585 7177
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   /sex="Male".
/lab host="B".
/lab host="B".
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
                                                                                /mol_type="genomic DNA"
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Laboratory Mouse DNA Resource
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CF339413
AJ661013
AL038582
AL038639
AL038839
CF330439
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AZ0525662
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TA35C12Q
TA380A07P
AG194579
CF334610
CL680297
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AZ774536
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ACCESSION: AL038839
ACCESSION: CF330439
ACCESSION: CN763587
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ACCESSION: AZ625662
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ACCESSION: AL038582
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ACCESSION:AL038460
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ACCESSION: AL454256
ACCESSION: AL497713
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           Jackson
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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KEYWORDS
SOURCE
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AZ486788/c
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                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0315 row: I column: 22
Seg primer: CGTTGTAAAACGACGGCCAGT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 24)
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Location/Qualifiers
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Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Utah
University of Utah
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from M. muscullus C57BL/6J (male) was obtained from the Jackso Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                     Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                         /mol_type="genomic DNA"
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http://www.jax.org/resources/documents/dnares/). The DNA

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REFERENCE
AUTHORS
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1 (bases 1 to 24)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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1M0429G03R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
Clone UUGC1M0429G03 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0429 row: G column: 03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
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                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C578L/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                      /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
(http://www.jax.org/resources/documents/dnares/). The DN:
was hydrodynamically sheared by repeated passage through
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0; Mismatches
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Mouse whole genome scaffolding with paired end reads from 10kb mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                       High quality sequence stop: 25.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0056 row: P column: 09
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ330737
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                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse 10kb plasmid UUGCIM library"./note="Vector: PW942nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                         /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                     clone="UUGC1M0056F09"
                                                                                                                                                                                                                                                                                        /db xref="taxon:10090"
                                                                                                                                                                                                                                 'sex="Male"
                                                                                                                                                                                                                                                                                                                                                                organism="Mus musculus"
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Pred. No. 11;
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KEYWORDS VERSION

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RESULT 5
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1520 AAAAAAAAAAGTAAAAGGG 1540
                                                                                                                                                                                                                                                                                                                                                                                     Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTAt 10.1) was mechanically shear to give a tight size distribution (4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
                                                                                                                                                                                                                                                                                          Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 18A, E-mail: barrell@eanger.ac.uk and nhl@eanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Trypanosoma brucei
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AL480935.1 GI:11846704
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                                                                                                                                                                                               /mol_type="genomic DNA"
/strain="TREU927"
                                                                                                                                                  /clone="231e08"
                                                                                                                                                                                                                                               organism="Trypanosoma brucei"
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Pred. No. 13;
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Pred. No. 11;
                                                    Mismatches
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AJ663467
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Best Local &
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Roslin, Midlochian, EH25 9PS, UNITED KINGDOM
Single pass sequencing. Bases called and trimmed with phred
v0.020425.c. Vector identified by cross match with the -minscore 20
and -minmatch 12 options. Vector:pBlueScriptII(KS+) R. Site 1:
ECORIR. Site 2: NotI Description: Normalised library constructed
from pooled tissue from day 30 placentas. Clones available from UK
Centre for Functional Genomics in Farm Animals, Roslin Institite,
Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.
                  Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
                                                                                                       Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                               Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (Dases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                               sequence.
CF291636
CF291636.1 GI:33660669
                                                                                                                                                                                                                                                                                                                                                                                                                            CF291636 RNA linear EST 14-AUG-2-14ROOT--02-C09.gl Rice root plasmid cDNA library (14ROOT) Oryzasativa (japonica cultivar-group) cDNA clone 14ROOT--02-C09, mRNA
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Contact: Anderson SI
Genomics and Bioinformatics
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Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sus scrofa
                                                                                       Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sus scrofa (pig)
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Kyeonggi, K
31 330 6193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone="C0000027_007"
/tissue_type="placenta"
/tissue_type="placenta"
/clone_Tib="CSEQRAN09"
/clone_Tib="CSEQRAN09"
/note="Vector: pBlueScriptII(KS+); Site_1: EcoRI; Site_2:
NotI; Single pass sequencing. Normalised library
constructed from pooled tissue from day 30 placentas."
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/db_xref="taxon:9823"
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Pred. No. 15;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AUTHORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM
                                                                                                                                                                                                                                                                                 source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            424 GTGGCGGCTGCGGCGCGACG 447
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                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0016 row: B column: 10
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, B., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AZ309553.1 GI:10350837 GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ309553 24 bp DNA linear GSS 29-SEP-20010016B10F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0016B10 F, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              308,
                                                                                                                                                                                                                                                                                                         quality sequence stop: 24. Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:39947"
/db_xref="taxon:39947"
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/dev_stage="14 days after germination"
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/clone_lib="Rice_root_plasmid_cDNA_library_(14ROOT)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA_was capped
with_oligoribonucleotides and then used as templates for
RT-pCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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                 /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PMD42nv, Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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                                                                                                                                                             'mol type="genomic DNA"
'strain="C57BL/6J"
'db xref="taxon:10000"
'clone="UUGC1M0016B10"
http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                      organism="Mus musculus"
                                                                                                                                            sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.3%;
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Pred. No. 16;
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TITLE JOURNAL COMMENT

REFERENCE

AUTHORS

FEATURES

source

Query Match Best Local

Similarity 83.3%;

Score 17.6; D Pred. No. 17; 0; Mismatches

DB 1;

Length 25;

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Indels

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Gaps

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Conservative

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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Cryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae; Streptophyta; Embryophyt
Spermatophyta; Magnoliophyta; Liliopsida; Poales;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25 pp mRNA linear EST 15-AUG-2003 HD--06-I14.gl OsHDAC1-overexpressing transgenic rice plasmid cDNA library (HD) Oryza sativa (japonica cultivar-group) cDNA clone HD--06-I14, mRNA sequence.
CF317007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193

Fax: 82 31 321 6355

Frail blacksons in the backsons of the Faxe of the Company of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Large-scale Sequencing Analysis
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CF317007.1 GI:33688768
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
                                             /note="Vector: pCR4-TOPO; Site_1: EcoRI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was
                                                                                                                                                                                                           /dev stage="proliferated callus on 2N6 media for 2 weeks" /lab_host="E.coli DH10B" /clone lib="OsHDAC1-overexpressing transgenic rice plasmid cDNA library (HD)"
derived from rice Histone Deacetylase overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was hydrodynamically sheared by repeated passage through
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                        tissue_type="callus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:39947"
/clone="HD--06-I14"
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Pred. No. 16;
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ACCESSION VERSION KEYWORDS

SOURCE

ORGANISM

RESULT 9 CF317007/c LOCUS

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                  RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seq primer: -21ml3
High quality seque
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Generation and analysis of 280,000 human expressed sequence tags
Genome Res. 6 (9), 807-828 (1996)
97044478
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        749097 25 bp mRNA linear EST 06-PEB-197008h08.sl Stratagene placenta (#937225) Homos sapiens cDNA clone IMAGE:70623 3' similar to similar to gb:862744 CLASS II HISTOCOMPATIBILITY ANTIGEN, M ALPHA CHAIN (HUMAN), mRNA sequence.
  AZ307896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      High qality sequence starts: 1
High qality sequence stops: 1
Source: IMAGE Consortium, LINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
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Contact: Wilson RK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: est@watson.wustl.edu
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                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                              /mol_type="nuc.../db_xref="GDB:491520"
/db_xref="H=xon:9606"
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/clone="IMAGE:70623"
                                                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Homo sapiens"
mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                         sex="male"
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RESULT 12
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                                                                                                                                                                1520 AAAAAAAAAAAAGTAAAAGGGA 1541
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Errr
Plate: 0010 row: N column: 11
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University of Utah Genome Center
University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T.,
Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
Niederhausern,A. and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1M0010N18F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0010N18 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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                                                                                                                                                                                                                                                                                              was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt end-sin high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwp42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                               Conservative
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'strain="C57BL/6J"
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                                                                                                                                                                                                                                                         Score 17.2;
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DEFINITION

CF308058 23 bp mRNA linear EST 15-AUG-ABF--01-L15.gl ABF3-overexpressing transgenic rice plasmid cDNA

EST 15-AUG-2003

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JOURNAL COMMENT
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AUTHORS
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb Nouse whole genome scaffolding with paired end reads from 10kb
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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24 bp DNA linear GSS 13-DEC-200
1M0474N20F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0474N20 F, genomic survey sequence.
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                     plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
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CF308058
CF308058.1 GI:33679819
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Location/Qualifiers
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Pred. No. 20;
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RESULT 14
CF300172/c
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VERSION
                                                                                                                              COMMENT
                                                                                                                                                                                                                                                     REFERENCE
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                                                                                                                                                   TITLE
JOURNAL
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                                                                                                                                                                                                                                                                       Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyl
Spermatophyta; Magnoliophyta; Liliopsida; Poales;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CF300172 23 bp mRNA linear EST 15-AUG-20
7LEAF--04-H15.b1 Rice leaf plasmid cDNA library II (7LEAF) Oryza
sativa (japonica cultivar-group) cDNA clone 7LEAF--04-H15, mRNA
                                                                                                                                                Kim, J.S., Jun, K.M., Cheong, P.J., Song, S.I., Kim, J.K., Kim, Y.-K. ar Large-scale Sequencing Analysis c Unpublished (2003)
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Insert Length: 1000 Std Err
Plate: 0474 row: N Column: 2
Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                         Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                CF300172.1 GI:33671933
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNDA2 (gi|473214|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and selected for ampicillin resistance."
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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s of Rice ESTs
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ACCESSION
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AL038576
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AUTHORS
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Matches 18
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Best Local Similarity
Matches 17; Conserv
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DKFZp566D1846 rl 566 (synonym: hfkd2) Homo
DKFZp566D1846, mRNA sequence.
AL038507
            AL038576 20 bj
DKFZp566F0346 rl 566 (synonym:
DKFZp566F0346, mRNA sequence.
AL038576
                                                                                                                     μ
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Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 20)
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AL038576.1
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   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                  Contact: MIPS
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ilarity 90.0%;
Conservative
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                                                                                                                                                                                                                    /organism="Homo sapiens"
/mol type="mRNA"
/db xref="taxoh:9606"
/clome="DKRZD56611846"
/tissue_type="kidney"
/tissue_type="kidney"
/dev_stage="fetal"
/lab_host="x1-Dblue"
/clome_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI; Site_2: Sa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
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/cb_xref="Nackdong"
/clone="TLEAF--04-H15"
/clone="TLEAF--04-H15"
/tissue_type="leaf"
/tissue_type="leaf"
/dev_stage="7 days after germination"
/lab_host="E.coli DH10B"
/clone lib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
with oligoribonucleotides and then used as templates for
RT-PCR."
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|mol_type="mRNA"
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GI:49682161
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Pred. No.
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Pred. No. 2;
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AL038754
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Best Local S
Matches 18
                                                   Matches
                                                                Query Match
Best Local
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               1518 TTAAAAAAAAAAAAGTAAAA 1537
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EST.
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                                                                                                                                                                                                                                                                              Contact: MIPS
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                                                                  Similarity
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L8; Conservative
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Unpublished (1999)
Contact: MIPS
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                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1. (bases 1 to 20)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Ottenwaelder, B., Obermaier, B.,
                                                                                                                                                                                                                                                           Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany.
Location/Qualifiers
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   Location/Qualifiers
                                                                                                                                             /organism="Homo sapiens"
/mol type="mRNA"
/db xref="texcon:9606"
/clone="DKFZp566M1346"
/tissue_type="kidney"
                                                                 /dev stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pamp1; Site_1: NotI; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="kidney"
/dev stage="fetal"
/lab_host="Xl-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: Not
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="DKFZp566F0346"
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90.0%; Pred. No. 22;
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0; Mismatches
                 Score 16.8;
Pred. No. 2:
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n: hfkd2) Homo
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                                                                                                                                                                                                                                                                                                                                                                                           Mewes,H.W.,
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                     22
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                                    DB 1;
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il Similarity 90.0%;
18; Conservation
                                                                                                                                                                                                                   Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 20)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CF298207

CF298207

TLEAF--01-H23.gl Rice leaf plasmid cDNA library II (7LEAF) Oryza sativa (japonica cultivar-group) cDNA clone 7LEAF--01-H23, mRNA
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1 (Dases 1 to 20)

Ottenwaelder, B., Obermaier, B., Mewes, H.W., Gassenhuber, J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DKFZp566P1746 rl 566 (synonym: DKFZp566P1746, mRNA sequence.
                                                                                                 Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EST (Ottenwaelder, et al.)
Unpublished (1999)
                                                                                                                                                             Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                CF298207.1 GI:33669968
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Location/Qualifiers
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                                                                                                                                                                                                    Contact: Nahm B.H.
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                                                            bhnahm@ggbio.com, bhna
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol type="mRNA"
/db_xref="taxon:9606"
/clone="bkrzp566p1746"
/tissue_type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: Not1; Site_2: Sa
/organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
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Pred. No. 22
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                                                                              bhnahm@bio.myongji.ac.kr
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>nym: hfkd2) Homo sapiens cDNA clone
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                                                                                                                                                                                                                                                                                                                                                                                                                                     source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0150 row: P column: 21
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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2M0150P21R Mouse 10kb plasmid
clone UUGC2M0150P21 R, genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Welss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1. (bases 1 to 20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 20.
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University of Utah
Rm. 308, Biomedical Polymers Rese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
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Similarity 90.0%;
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /dev_stage="7 days after germination"
/lab_host="E.coli DHLOB"
/clome_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/clome_Tib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site 1: ECCRI; mRNA was capped
with_oligoribonucleotides and then used as templates for
                                                                                                                                                                   Laboratory Mouse DNA Resource
                                                                                                                                                                                            /lab host="E. Coli strain XL10-Gold, Tl-resistant, F-"/clome_lb="Mouse 10kb plasmid UUGCIM library" /note="Vector: PW927v; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jacksor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RT-PCR."
                                                                                                                                                                                                                                                                                                                                                             /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
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/clone="7LEAF--01-H23"
/tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                            db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                             'sex="Male"
                                                                                                                                                                                                                                                                                                                   clone="UUGC2M0150P21"
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Pred. No. 22;
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smid UUGC1M library Mus musculus genomic
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                                                                                                                                                                                                 Jackson
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2M0203K21R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0203K21 R, genomic survey sequence.
AZ943299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0203 row: K column: 21
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2000)
Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ943299.1 GI:13807290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTAAAAAAAAAAAAAAAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pw042 (gi|4732.14|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                              /lab_host="E. coli strain XL10-Gold, T1-resistant, F/clone_lib="Mouse 10kb plasmid UUGC2M library" /note="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (female) was obtained from the Jamusculus C57BL/6J (female)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0203K21"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                     'sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.2%;
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Pred. No. 22
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COMMENT

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RESULT 22
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KEYWORDS
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                        CF310486 22 bp mRNA linear EST 15-AUG-2003 ABF--05-C16.gl ABF3-overexpressing transgenic rice plasmid cDNA library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone ABF--05-C16, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AL038477

DKFZp566C1646_r1 566 (synonym:
                                                                                  EST.
                                                                                                     CF310486
CF310486.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 22)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AL038477.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: MIPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EST (Ottenwaelder, et al.)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ottenwaelder, B., Obermaier, B., Mewes, H.W., Gassenhuber, J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           electrophoresis. Vector DNA was prepared from a derivative of pWp42 (gi|4732114|gb|AFL19072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: paMP1; Site_1: NotI; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="DKFZp566C1646"
/tissue_type="kidney"
/dev_stage="fetal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="mRNA"
/db_xref="taxon:9606"
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                                                                                                       GI:33682247
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90.0%; Pred. No. 25;
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 Mismatches

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Pred. No. 24
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hfkd2) Homo sapiens
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FEATURES

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SOURCE

FOCUS

CF310486

SOURCE

Ehrhartoideae; Oryzeae; Oryza

KEYWORDS

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RESULT 24
CF329694/c
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KEYWORDS
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AUTHORS
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JOURNAL
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Matches
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                                                                                                                                                                                                                                               Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
                                                                                                                                                                                                                                                                                                             1 (bases 1 to 23)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae,
Ehrhartoideae, Oryzeae, Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CF329694 23 bp mRNA linear EST 18-AUG-2003
NACL--05-B19.b1 Rice callus plasmid cDNA library (NACL) Oryza
sativa (japonica cultivar-group) cDNA clone NACL--05-B19, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 22)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
Contact: Nahm B.H.
                                                                                                                                                                                                   Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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EST.
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                                                                                                                                                          bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Vector: pCR4-TOPO; Site_1: EcoRI; Leaf was dried for Ahrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /dev stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone_lib="ABF3-overexpressing transgenic rice
cDNA library (ABF)"
/organism="Oryza sativa (japonica cultivar-group)"
/mol type="mRNA"
/culTivar="Nackdong"
/db xref="taxon:39947"
/clone="NACL--05-B19"
/tissue_type="callus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Oryza sativa
/mol_type="mRNA"
/cultivar="Nackdong"
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/clone="ABF--05-C16"
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Pred. No. 25;
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DEFINITION

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C0000045_P10, mRNA

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AUTHORS
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CF333801/c
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                                        AJ668301/c
LOCUS
                                                                              RESULT 26
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AJ668301
AJ668301
sequence.
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JMT--02-N11.gl AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT) Oryza sativa (japonica cultivar-group) cDNA clone JMT--02-N11, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech I.
Genomics and Bioinformatics, MyongJi University
Yongin, Kyeenggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CF333801
CF333801.1 GI:33815910
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                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                             /note="Vector: pCR4-TOPO; Site_1: EcoRI; Oligo-capped mRNA
was reverse transcribed and then used for PCR. mRNA was
pepared from Arabidopsis Jasmoinate Carboxyl
                                                                                                                                                                                                                                                                                                                                                                    /dev stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone lib="ActMT-overexpressing transgenic rice plasmid
cDNA library (JMT)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone lib="Rice callus plasmid cDNA library (NACL)" /note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped with oligoribonuclectides and then used as templates for
                                                                                                                                                                                                                                                                                           methyltransferase overexpression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                tissue_type="leaf"
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'clone="JMT~-02-N11"
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mol_type="mRNA"
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lab_host="B.coli DH10B"
                                                                                                                                                                                                                 1.2%; Score 16.8; 1
90.0%; Pred. No. 26;
tive 0; Mismatches
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Pred. No. 26;
0; Mismatches
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RESULT 27
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Best Local
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                                                                                                                                                         response genes
Genome Biol. 4
22881942
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Sus scrofa
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Glossina morsitans morsitans
Glossina morsitans morsitans
Elsaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Hippoboscoidea; Glossinidae; Glossina.
1 (bases 1 to 23)
Tehane M.T. Alfoy C. Gibcon W. Kerbornov B. Berriman M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23 bp mRNA linear EST 10-C EX550903 Glossina morsitans morsitans adult infected gut Glo morsitans morsitans morsitans morsitans cDNA clone Tsell5e01_qlc, mRNA sequence. EX550903
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Anderson, S.I., Finlayson, H.A. and Archibald, A.L.
Development of cDNA and EST resources for studying reproduction embryo development in pigs and cattle
Unpublished (2004)
                               Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
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BX550903.1 GI:33374645
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Genomics and Bioinformatics
                                                                                                                    14519198
Contact: Hall N
                                                                                                                                                                                                                 morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                           Lehane,M.J., Aksoy,S., Gibson,W., Kerhornou,A., Bemilton,J., Soares,M.B., Bonaldo,M.F., Lehane,S.
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Mammalia; E
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AJ668301.1
                                                                                                                                                                                                                                        Adult midgut expressed sequence tags from the tsetse fly Glossina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                    M.J.Lehane
of Biological Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  constructed
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/mol_type="mRNA"

/db_xref="taxon:9823"
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Pred. No. 2
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3. and Hall,
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Sus.
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REFERENCE
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Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Hippoboscoidea; Glossinidae; Glossina.
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morsitans morsitans cDNA clone Tse91f03_p1c, mRNA sequence.
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Bangor LL57 2UW
All clones with suffix q1c are reverse
end of the cDNA all p1c reads are from
the 3' end.
                                                                                                                                                                                                                                                                                                                          Prof. M.J.Lehane
School of Biological Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        response genes
Genome Biol. 4 (10), R63 (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lehane,M.J., Aksoy,S., Gibson,W., Kerhornou,A., Berriman,M., Hamilton,J., Soares,M.B., Bonaldo,M.F., Lehane,S. and Hall,N. Adult midgut expressed sequence tags from the tsetse fly Glossina morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                       All clones with suffix qlc are revend of the cDNA all plc reads are the 3' end.
                                                                                                                                                                                                                                                                                  University of Wales,
Bangor LL57 2UW
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                                 /clone="Tse91f03_p1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone="Tsel15e01 q1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Glossina morsitans morsitans"
/mol_type="mRNA"
/mub_species="morsitans"
                                                                                                                   sub_species="morsitans"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note="country: Zimbabwe; EST from adult gut infected with
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                                                                                                                                      mol_type="mRNA"
                                                                                                                                                           organism="Glossina morsitans morsitans"
                                                                                                                                                                                                  ocation/Qualifiers
note="country: Zimbabwe; EST from adult gut infected with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GI:33434952
                                                                                                  xref="taxon:37546"
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AZ486853/c
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Best Local Similarity
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AZ486853.1 GI:10654033
GSS.
Mus musculus (house mouse
Mus musculus
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0315 row: M column: 14
Seq primer: CACACAGGAAACAGCTATGACC
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Fax: 801 585 7177
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Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse whole genome scaffolding with paired end reads from 10kb
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1M0315M14R Mouse 10kb plasmid UUGC1M library Muclone UUGC1M0315M14 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B.
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/lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PMD42nv; Purified genomic DNA from M.
musculus C57BL/GJ (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blint end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pwm92 (gil4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Cold (Stratagene) cells
and selected for ampicillin resistance."
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strain="C57BL/6J"
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Pred. No. 30;
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1M0510005R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC1M0510005 R, genomic survey sequence.
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University of Utah Genome Center
University of Utah
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Location/Qualifiers
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Insert Length: 10000 Std Error: (
Plate: 0510 row: O column: 05
Seq primer: CACACAGGAAACAGCTATGACC
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Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von
Niederhausern, A. and Wright, D., Weiss, R.
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Dunn, D., Aoyagi, A., Barber, M.,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tel: 801 585 5606
Fax: 801 585 7177
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                     was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymicleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi 4732114 |gb|AP129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: pWD42TN', Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
and selected
                                                                                                                                                                                                                                                                                                                                                                                                                   Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                        (http://www.jax.org/resources/documents/dnares/). The DNA
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Pred. No. 30;
  for ampicillin resistance."
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ORGANISM
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1M0529F08F Mouse 10kb plasmid UUGC1M 11D1a17 ....
-1-ne UUGC1M0529F08 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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Mammalia; Eutheria; Rodentia;
1 (bases 1 to 19)
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GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0529 row: F column: 08
Seq primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2000)
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Location/Qualifiers
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10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 60.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                               /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biomedical Polymers Research Bldg.,
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/strain="C57BL/6J"
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Query Match Best Local Similarity

1.2%;

Score 16.4; Pred. No. 27

DB 1;

Length 19

Matches Query Match Best Local (

Similarity

94.4%;

Score 16.4; D. Pred. No. 29; 0; Mismatches

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AUTHORS
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Insert Length: 10000 Std Error:
Plate: 0077 row: E column: 20
Seq primer: CGTTGTAAAACGACGGCCAGT
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1M0077E20F Mouse 10kb plasmid
clone UUGC1M0077E20 F, genomic
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2000)
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNB42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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RESULT 34
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T. brucei sheared genomic Di
genomic survey sequence.
AL497383 1 GI:11865504
GSS.
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Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for Clones, please contact: Mike Lehane
Trypanosoma brucei
Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lehane, M.J., Aksoy, S., Gibson, W., Kerhornou, A., Berriman, M., Hamilton, J., Soares, M.B., Bonaldo, M.F., Lehane, S. and Hall, N. Adult midgut expressed sequence tags from the tsetse fly Glossina morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Hippoboscoidea; Glossinidae; Glossina.
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Bangor LL57 2UW
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School of Biological Sciences,
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of the cDNA all plc reads are from
3' end.
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                                                                                                                                                                                                                                                                                                                                                              /note="country: Zimbabwe;
T.brucei"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Glossina morsitans morsitans"
/mol type="mRNA"
/sub_species="morsitans"
                                                                                                                                                                                                                                                                                                                                                                                                             /clone="Tse101g03_p1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans mo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                /sub_species="morsitans"
/db_xref="taxon:37546"
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                                       Prof. M.J.Lehane
School of Biological Sciences,
University of Wales,
Bangor LL57 2UW
All clones ....
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BX564412 Glossina morsitans morsitans adult infected gut Glossina morsitans morsitans cDNA clone Tse7le10_plc, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17;
All clones with suffix qlc are reverse primer reads starting at end of the cDNA all plc reads are from the 3' end.
                                                                                                                                                                                                                                                                                                                                 response genes
Genome Biol. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Hippoboscoidea; Glossinidae; Glossina.
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Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      to give a tight size distribution (
4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
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Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
                                                                                                                                                             Hinxton, Cambridge, CB10 1SA, UK Request for clones, please contact:
                                                                                                                                                                                                                                                           Contact: Hall N
                                                                                                                                                                                                                                                                                       14519198
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/strain="TREU927"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /db_xref="taxon:5691"
/clone="303g05"
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Pred. No. 33;
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                                                                                                                                                                    Mike Lehane
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0012 row: H column: 15
Seg primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb Nouse whole genome scaffolding with paired end reads from 10kb
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AZ308846.1 GI:10349246
GSS.
Mus musculus (house mous
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              University of Utah Genome Center University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                  quality sequence stop: 21.
Location/Qualifiers
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                   (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
                                                                                                                                 /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
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/tissue type="adult infected gut"
/clone_Tib="Glossina morsitans morsitans adult infected
polymucleotide kinase. Adaptor oligonucleotides
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                                                                                                                     Laboratory Mouse DNA Resource
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/db_xref="taxon:37546"
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                                                                                                                                                                                                                                     sex="Male"
                                                                                                                                                                                                                                                           clone="UUGC1M0012H15"
                                                                                                                                                                                                                                                                                    db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                       /mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                        organism="Mus musculus"
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Pred. No. 35;
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AZ493766
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ORGANISM
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1M0328C11R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0328C11 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0328 row: C column: 11
Seq primer: CACACAGGAAACAGCTATGACC
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 21)
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High quality sequence stop: 21.
Location/Qualifiers
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                         /organism="Mus musculus"
/mol type="genomic DNA"
/strain="C57BL/6J"
/db xref="taxon:10090"
/clone="UUGC1M0328C11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWH92 (gi/4732114/gb/AF129072.1), a copy-number inducible derivative of plasmid RI. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA (http://www.jax.org/resources/documents/dnares/) through a was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymerecide kinase. Adaptor oligonucleotides were polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                    /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse 10kb plasmid UUGCIM library/notes-Tvetor: PWPA2ny; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
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Pred. No. 35;
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AZ513902/c
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Best Local
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb Teachtliched (2004)
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Class: plasmid ends
High quality sequence stop: 21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0360 row: A column: 13
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Fax: 801 585 7177
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
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GSS.
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1M0360A13F Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC1M0360A13 F, genomic survey sequence.
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv, Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                   db_xref="taxon:10090"
clone="UUGC1M0360A13"
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Pred. No. 3:
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AZ626965
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Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 21)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Tislam,H., Rose,R., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1M0467E15R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0467E15 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                University of Utah Genome Center University of Utah
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Class: plasmid ends
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Plate: 0467 row: B column: 15
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Fax: 801 585 7177
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel
                                                                                                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /Clone lib="mouse 10kb plasmid UUGCIM library" /note="Vector: PW042nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Biomedical Polymers Research Bldg., 20 S.
                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                         /clone="UUGC1M0467E15"
                                                                                                                                                                                                                                                                                                                                                                                                    db xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                 'sex="Male"
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Pred. No. 35;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Trace considered overall poor quality
Insert Length: 520 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 1.
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AA911173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 22)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
cound through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGTTGTGTGTGTGTGCCTG 21
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Conservative
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                                                                                           /clone lib="NCI CGAP_Kid3"
/clone lib="NCI CGAP_Kid3"
/notes Torgan: kidney; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Ecc RI; 1st strand cDNA was primed with a Not I - oligo(dT) primer, double-stranded cDNA was ligated to Ecc RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Ecc RI sites of the modified pT7T3 vector. mRNA source: 2 pooled kidneys. Library went through one round of normalization. Library constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:9606"
/clone="IMAGE:1520346"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Homo sapiens"
|mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ocation/Qualifiers
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                    1.2%;
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                    Score 16.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0542 row: D column: 04
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Unpublished (2000)
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1M0542D04F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
Clone UUGC1M0542D04 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Class: plasmid ends
High quality sequence stop: 23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              308,
was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of phasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone_lib="Mouse 10kb plasmid UUGCIM library"/notee="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="WGC1M0542D04"
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/strain="C57BL/6J"
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Best Loc Matches

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1517 ATTAAAAAAAAAAAAAGTAAAA 1537
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                 l Similarity
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seq primer: CGTTGTAAAACGACGGCCAGT Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0059 row: J column: 16
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Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
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                                                                              1.2%;
llarity 85.7%;
Conservative
                                                                                                                                                                        Imboratory Mouse DNA Resources

(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gil 4732114 | gb| AFT29072 1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/clone_lib="Mouse lokb plasmid UUGCIM library"
/note="Vector: pwp2zpv; burified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
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SOURCE
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        Query Match
Best Local Similarity
Matches 16; Conserv
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2M0222H16R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0222H16 R, genomic survey sections
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Location/Qualifiers
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Insert Length: 10000 Std Error: (
Plate: 0222 row: H column: 16
Seq primer: CACACAGGAAACAGCTATGACC
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus
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Fax: 801 585 7177
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84112, USA
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           Conservative
                                                                                                       adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /lab host="E. coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGC2M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
/musculus C57BL/6J (female) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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/strain="C57BL/6J"
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100.0%; Pr
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     Score 16; DB 1; Length 21; pred. No. 40; 0; Mismatches 0; Indels
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AZ345795
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17; Conserv
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University of Utah Genome Center
University of Utah
University of Utah
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Dunn, D., Aoysgi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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Insert Length: 10000 Std Error: 0.00
Plate: 0080 row: H column: 09
Seg primer: CACACAGGAAACAGCTATGACC
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High quality sequence stop: 19.
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Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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AZ345795.1 GI:10425032
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                                                                                                                                                                                                                                                                          (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA.
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                                                                                                                                                                                                         adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   db_xref="taxon:10090"
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                                                                                                                             1.1%;
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                                                                                                                                Score 15.8;
Pred. No. 41
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mid UUGC1M library Mus musculus genomic
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Nouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads from 10kb
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Insert Length: 10000 Std Error: 0.00
Plate: 0520 row: P column: 13
Seq primer: CACACAGGAAACAGCTATGACC
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Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Location/Qualifiers
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Fax: 801 585 7177
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                                                                         was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was bint end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt end-sified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi 4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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1.1%;
   Score 15.8;
Pred. No. 41;
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smid UUGC1M library Mus musculus genomic
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                              Length 19;
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Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                                                                                                                     DKFZp566A1946_r1 566 (synonym: DKFZp566A1946, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 20)
Ottenwaelder, B., Obermaier, B., Mewes, H.W.,
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                                                                                                                                                                                                                                                                            Ottenwaelder, B., Obermaier, B.,
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Location/Qualifiers
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Unpublished (1999)
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Mammalia; Eutheria; Primates;
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Location/Qualifiers
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/mol types"mRNA"
/mol types"mRNA"
/db xref="taxon:9606"
/clone="DKK72p566A1946"
/tlssue_types"kidney"
/dev_stage="fetal"
/lab hoste"X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI; Si
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/clone_lib="566 (8ynonym: hfkd2)"
/note="Vector: pAMP1; Site_1: Not:
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/tissue_type="kidney"
/dev_stage="fetal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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Pred. No. 43
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nym: hfkd2) Homo
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hfkd2) Homo sapiens cDNA clone
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AL587759/c
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Dept. Genom:
Roslin Institute
Roslin, Midlothian, EP
Tel: +44 (0)131 527 4;
Tex: +44 (0)131 440 0
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                                                                                                                                                                                Gallus gallus
                                                                                                                                      Eukaryota; Metazoa; Chordata; (Archosauria; Aves; Neognathae; Phasianinae; Gallus.
                                                                                                                                                                                                                                           AL587759 20 bp
AL587759 BP Chicken Brain Library
ROS061G06, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DKFZp566F1746, mRNA sequence.
                                                                           Contact: Frazer Murray
                                                                                        Unpublished (2001
                                                                                                   Murray, F.
BP Chicken Brain Library
                                                                                                                                                                                            Gallus gallus (chicken)
                                                                                                                                                                                                                      AL587759.1 GI:13192793
                                                                                                                                                                                                                                     AL587759
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Location/Qualifiers
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Unpublished (1999)
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Ottenwaelder, B., Obermaier, B.,
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Mammalia; Eutheria; Primates;
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                                                              Genomics and Bioinformatics
                                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="kidney"
/dev stage="fetal"
/lab_host="X1-Zblue"
/clome_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="DKFZp566F1746"
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Pred. No. 4:
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Pred. No. 43;
                                     9PS, UK
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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AZ345856.1 GI:10425093
GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.00
Plate: 0080 row: G column: 17
Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ345856

1M0080G17R Mouse 10kb plasmid clone UUGC1M0080G17 R, genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   84112, USA
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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GCGGCCGTTTTTTTTTTTTTTTTT 3' Poly A RNA purchased from Clonetech
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 308,
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Location/Qualifiers
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/note="Vector: pSPORT1; Site 1: Not1; Site 2: Sal1; Cloned
/note="Vector: pSPORT1; Site 1: Not1; Site 2: Sal1; Cloned
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/note="Vector: pSPORT1; Site 1: Not1; Site 2: Sal1; Sit
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/dev_stage="Unknown"
/lab_host="DH10B"
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F
/clome_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                'sex="Male"
                                                                                                                                                 clone="UUGC1M0080G17"
                                                                                                                                                                               db_xref="taxon:10090"
                                                                                                                                                                                                             /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                        organism="Mus musculus"
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'mol_type="mRNA"
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Pred. No. 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  genomic survey sequence.
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UUGC1M library Mus musculus
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SOURCE
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AZ486784/c
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weise,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ486784 20 bp DNA linear GSS 0:
1M0315C20F Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC1M0315C20 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                  Seq primer: CGTTGTAAAACGACGGCCAGT Class: plasmid ends High quality sequence stop: 20.
                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0315 row: C column: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducable derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                             /db_xref="taxon:10090"
/clone="UUGC1M0315C20"
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
/note="Wecttor: PWB42nv; Burified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                    /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                        organism="Mus musculus"
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                                                                                                                                                                                                                                                                                         Location/Qualifiers
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Pred. No. 43
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ACCESSION VERSION

KEYWORDS

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Matches

REFERENCE AUTHORS

COMMENT

JOURNAL

TITLE

FEATURES

Laboratory Mouse DNA Resource

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REFERENCE
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VERSION
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AZ858419
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                  source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 20)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R. Mouse, whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0163 row: O column: 03
Seq primer: CACACGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 20.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plasmid inserts
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F./clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biomedical Polymers Research Bldg., 20 S. 2030 E.,
                                                                                                                                                              db_xref="taxon:10090"
clone="UUGC2M0163003"
                                                                                                                                                                                                             /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                         organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                              ocation/Qualifiers
                                                                                                                                       sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.1%;
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Pred. No. 43;
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REFERENCE AUTHORS TITLE JOURNAL

COMMENT

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ACCESSION

SOURCE

ORGANISM

LOCUS DEFINITION RESULT 53 AL048772 밁 S

RESULT 54 CF318152/c LOCUS DEFINITION

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Best Local
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CF318152 21 bp mRNA linear EHD--08-C11.bl OsHDAC1-overexpressing transgenic rice library (HD) Oryza sativa (japonica cultivar-group) cHD--08-C11, mRNA sequence.
                                                                                                                                                                                                                                                                                                    17;
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DKFZp566N143 rl 566 (synonym:
DKFZp566N143, mRNA sequence.
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1 (bases 1 to 21)

Koehrer, K., Beyer, A., Mewes, H.W., Gassenhuber, J. and Wiemann, S. EST (Koehrer, et al.)

Unpublished (1999)
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                       n 1.1%;
Similarity 89.5%;
                                                                                                                                                                                                          GTTAAAAAAAAAAAAAAAA 20
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                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:9606"
/clone="DKPZp566N143"
/tissue type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agazose gel electrophoresis. Vector DNA was prepared from a derivative of pwp82 (gi|4732114|gb|sp129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Homo sapiens"
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Pred. No. 4
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RESULT 55
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17; Conserv
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                                                                                                                                                                                                                                                                                                        Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; rosids; Vitaceae; Vitis.

1 (bases 1 to 21)
1 (bases 1 to 21)
1 (bases 2)
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EST.
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Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                               or seeds) at Various Developmental Stages
Unpublished (2002)
Contact: Hamdi S.
CMR 619 - Equipe Biologie de la Vigne
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CN546489.1 GI:46911114
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                                                                                                    Agronomique
                                                                                                                                     Universite de Bordeaux I,
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: 82 31 330 6193
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                                                               Avenue Edouard Bourleaux,
00-33-(0)5-57-12-25-50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /dev stage="proliferated callus on 2N6 media for 2 weeks" /lab_host="E.coli DH10B" /clone lib="OsHDAC1-overexpressing transgenic rice plasmid cDNA library (HD)"
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/clone="HD--08-C11"
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/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tissue_type="callus"
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Pred. No. 46
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                                                                      33883 Villenave D'Ornon Cedex,
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RESULT 56
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JOURNAL
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                                                                                                                                                                                                                                                                                                                  Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 21.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0436 row: G column: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 21)
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Unpublished (2000)
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Email: s.hamdi@bordeaux.inra.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert
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USA
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: pW042nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resourcee/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
o.005 inch orifice at constant velocity. The sheared DNA
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/clone_lib="Ripe Grape Berry Lambda Triplex2 Library"
/note="forgan: Fruit without seeds; Vector: Lambda
TriplEx2; Site_1: SfiIA; Site_2: SfiIB; Oriented library"
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/mol type="mRNA"
/culfivar="Cabernet Sauvignon"
/db xref="caxon:29760"
/clone="B3CS57RB007E11"
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                                                                                                                                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                           organism="Mus musculus"
                                                                                                                                                                      sex="Male"
                                                                                                                                                                                           clone="UUGC1M0436G12"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B. Weiss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.8; D)
Pred. No. 46;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                     Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mid UUGC1M library Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                genomic
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was blunt end-repaired with T4 DNA polymerase and T4

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RESULT 57
AZ764492
LOCUS
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AUTHORS
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ORGANISM
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VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus (house mouse)
Mus musculus
Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 21)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T.,
Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von
Niederhausern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ764492 21 bp DNA linear GSS 16-FEB-200
1M0560D04R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
Clone UUGC1M0560D04 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0560 row: D column: 04
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ764492.1 GI:12879511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                              lass: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pW042 (gi[4732114]gb[AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
                                                                                                    /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCM library"
/note="Vector: PMD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
                                                                                                                                                                                                                                                              /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0560D04"
                                                                                                                                                                                                                                                                                                                                                                  organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                     ocation/Qualifiers
                                                                                                                                                                                                                                        sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymers Research Bldg., 20 S. 2030 E.,
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Pred. No. 46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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COMMENT

Prof. M.J.Lehane School of Biological Sciences,

University of Wales, Bangor LL57 2UW

clones with suffix q1c are reverse primer reads starting
 of the cDNA all p1c reads are from
 3' end.

/clone="Tse24f09_p1c" /tissue_type="adult infected gut" /clone_lb="Glossina morsitans morsitans adult infected

note="country: Zimbabwe; EST from adult gut infected with

/organism="Glossina morsitans morsitans" |mol type="mRNA" |sub species="morsitans" |db_xref="taxon:37546"

JOURNAL MEDLINE PUBMED

FEATURES

ocation/Qualifiers

Bource

Query Match Best Local

Similarity

1.1%; Score 15.8; 89.5%; Pred. No. 48

0; Mismatches

2

0

0

48;

DB 1; Length 22;

.brucei "

REFERENCE AUTHORS

TITLE

VERSION KEYWORDS SOURCE

ORGANISM

ACCESSION

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LOCUS
DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 58
BX556059/c
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1518 TTAAAAAAAAAAAGTAAA 1536
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Glossina morsitans morsitans
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BX556059 22 bp mRNA linear EST 10-OCT-200 BX556059 Glossina morsitans morsitans adult infected gut Glossina morsitans morsitans cDNA clone Tse24f09_plc, mRNA sequence.

BX556059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 TTAAAAAAAAAAAAAAAA 21
                                                                                                                                                                                                                 response genes
Genome Biol. 4 (10), R63 (2003)
                                                                                                                                                                                                                                                                             Lehane, M.J., Aksoy, S., Gibson, W., Kerhornou, A., Berriman, M., Hamilton, J., Soares, M.B., Bonaldo, M.F., Lehane, S. and Hall, N. Adult midgut expressed sequence tags from the tsetse fly Glossina morsitans morsitans and expression analysis of putative immune
Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BX556059.1 GI:33380008
                                                                                                                      Contact: Hall N
                                                                                                                                                            14519198
                                                                                                                                                                                                                                                                                                                                                                                                                            Hippoboscoidea; Glossinidae; Glossina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Glossina morsitans morsitans
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ligated to the blunt ends in high modar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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ligated to the blunt ends in high molar excess. The
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89.5%;
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Pred. No. 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
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CO778290
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AUTHORS
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CF310806/c
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                                                                                                                               SOURCE
                                                                                                                                                                                                 ACCESSION
                                                                                                                                                                                                                                           DEFINITION
                                                                                                                                                                                                                                                                 FOCUS
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                             22
                                                                                                                                                                      CO778290 22 bp mRNA linear EST 05-AUG-2004 BL003B H01 6-Day Axolotl Tail Blastema (6DAxBL) Ambystoma mexicanum cDNA 57 similar to hypothetical protein, mRNA sequence.
CO778290 CO778290.1 GI:50994270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta;
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae;
Ehrhartoideae; Oryzeae, Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CF310806 22 bp mRNA linear EST 15-AUG-2003
ABF--05-K20.gl ABF3-overexpressing transgenic rice plasmid cDNA
library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 22)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                           Ambystoma mexicanum
Eukaryota; Metazoa; Chordata;
Amphibia; Batrachia; Caudata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CF310806
CF310806.1 GI:33682567
Habermann, B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Nahm B.H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABF--05-K20, mRNA sequence.
                                           Ambyвсоma.
                                                                                                                             Ambystoma mexicanum (axolotl)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TAAAAAAAAAAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAAAAGGG
                    (bases 1 to 22)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bhnahm@ggbio.com, bhn
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Vector: pCR4-TOPO; Site 1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="leaf"
/dev_stage="14 days after germination"
/lab_host="E.col1 DH10B"
/clone lib="ABF3-overexpressing transgenic rice plasmid cDNA library (ABF)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Oryza sativa (japonica cultivar-group)"
/mol type="mRNA"
/cultivar="Nackdong"
/db xref="taxon:39947"
/clone="ABF--05-K20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
Bebin, A.G.,
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Pred. No. 48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
Herklotz, S.,
                                                             Craniata; Vertebrata; Euteleostomi; Salamandroidea; Ambystomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
Volkmer, M., Eckelt, K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>ب</u>
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2M0098K07F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0098K07 F, genomic survey sequence.
AZ823875
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Contact: Elly
                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0098 row: K column: 07
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 22.
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Tel: 0049 351 210 2620
Fax: 0049 351 210 1489
Email: tanaka@mpi-cbg_de
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                                                                                                                                                                                                                    University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                             plasmid inserts
Unpublished (2000)
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Seq primer: GCA CAT TAG GCC TAT
Location/Qualifiers
                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                          Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                        Mouse whole genome scaffolding with paired end reads from 10kb
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/cell type="regenerating tail blastema"
/cell type="regenerating tail blastema"
/clone_lib="6-Day Axolot! Tail Blastema (6DAxBL)"
/note="Vector: pCMVSportt; Site_1: NotI; Site_2: Sall;
/normalized cDNA plasmid library prepared by Invitrogen.
Vinormalized cDNA plasmid library prepared by Invitrogen.
Size fractionated mRNA was polydT primed and cloned into
NotI-Sall site of pCMVSport6. Bacterial host is
EMDH10B-TONA. Average insert size is 1.67 kB.
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/mol_type="mRNA"
/db_xref="taxon:8296"
    Location/Qualifiers
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Pred. No. 48;
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BH000233
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                                                                                       Tel: 801 585 5606

Fax: 801 585 7177

Fmai: ddunn@genetics.utah.edu
Email: ddunnggenetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0287 row: L column: 21
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota, Metazoa; Chordata, Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 22)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BH000233 22 bp DNA linear GSS 27-APR-200200287L21R Mouse 10kb plasmid UUGC2M library Mus musculus genomic clone UUGC2M0287L21 R, genomic survey sequence
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                        High quality sequence stop: 22.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                    Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                      USA
                                                                                                                                                                                                                                                            Biomedical
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/note="Vector: PWD42nv; Purified genomic DNA f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DNA"
/strain="C57BL/6J"
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Best Local
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1520 AAAAAAAAAAAAGTAAAAG 1538
                                                                                                                                                                                                                                                                                    Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

(bases 1 to 22)

Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R. and Wilson,R. WashU-NCI human EST Project
Unpublished (1997)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AI708831 22 bp mRNA linear EST 04-JUN-1999 as27d12.x1 Barstead aorta HPLRB6 Homo sapiens cDNA clone IMAGE.2318423 3' similar to TR:Q33563 Q33563 EATRO 164 KINETOPLAST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 AAAAAAAAAAAAAAAAAAA 21
                                                                                                                                                                                                      Seq primer: -40UP from Gibco
High quality sequence stop: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mRNA sequence.
                                                                                                                                                            quality sequence stop: 1.
Location/Qualifiers
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314 286 1810
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/Clone lib="Mouse 10kb plasmid UUGC2M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (female) was obtained from the Jackson
Laboratory Mouse DNA Resource
Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
organism="Homo sapiens"
'mol_type="mRNA"
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/strain="C57BL/6J"
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Pred. No. 48;
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RESULT 65
AW332181/c
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AUTHORS
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VERSION
KEYWORDS
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AJ747407
   DEFINITION
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Best Local Similarity
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AW332181 22 bp mRNA linear EST S5C7 AGS-1 Pneumocystis carinii cDNA 3', mRNA sequence.
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AJ747407 forward - stimulated minus unstimulated macrophage Sus
scrofa cDNA clone F_C0001825b_A06, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Hopwood PA

Contact: Hopwood PA

Dept. of Preclinical Veterinary Sciences

Royal School for Veterinary Studies

Summerhall, Edinburgh, EH9 1QH, UNITED KINGDOM

Sequencing was performed by ARK genomics. This clone is available

Sequencing was performed by ARK genomics. Midlothian EH25 9PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Development of a porcine cDNA microarray Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Euthería; Cetartiodactyla; Suina; Suidae; Sus.

1 (Dases 1 to 22)

Hogwood, P.A., Zhang, F., Lowden, S., Talbot, R., Burt, D., Archibald,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sus scrofa
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AJ747407.1 GI:49917653
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                                                                                                                                                                                                                                                                                                                                                                                                                     See www.ark- genomics.org or contact info@arkgenomics.org.
Location/Qualifiers
                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                          /mol_type="mRNA"
/db_xref="taxon:9823"
/clone="F_C0001825b_A06"
/tissue_type="lung"
/cell_type="macrophage"
/clone_lib="forward - stimulated minus unstimulated
                                                                                                                                                                                                                                                                                                                                                                                   organism="Sus scrofa"
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/clone="IMAGE:2318423"
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Pred. No. 54
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Pred. No. 54;
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                                                                                                                                                                                     Mismatches
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                    EST 31-JAN-2000
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SOURCE
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AW332399/c
LOCUS
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AW332399.1
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                                                                                                                                                                                                                                                                                                                                              Pneumocystis carinii
Eukaryota; Fungi; Ascomycota; Pneumocystidomycetes;
Pneumocystidaceae; Pneumocystis.
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SBA2 AGS-1 Pneumocystis carinii
מייימים
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101 Morgan Building, University of Kentucky, Lexington,
40506-0225, USA
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Smullan,A.G., Arnold,J., Weise,M., Wunderlich,J.,
Edman,J.C., Kovacs,J. and Cushion,M.
Expressed sequence tags from Pneumocystis carinii
Unpublished (2000)
                                                                                                                                                                  School of Biological Sciences
University of Kentucky
101 Morgan Building, Universi
40506-0225, USA
                                                                                                                                                                                                                                        Unpublished (2000)
Contact: Staben C
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Smulian,A.G., Arnold,J., Weise,M., Wunderlich,J.,
Edman,J.C., Kovacs,J. and Cushion,M.
Expressed sequence tags from Pneumocystis carinii
                                                                                                                        Tel: 606 257 2161
Fax: 606 257 1717
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Location/Qualifiers
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AW332181.1
                                                                                        Email: staben@pop.uky.edu.
Location/Qualifiers
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/Clone lib="AGS-1"
/note="Vector: Lambda ZAP II; Site_1: EcoRI; Site_2: XhoI;
/note="Vector: Lambda ZAP II; Site_1: EcoRI; Site_2-6,
P. carinii organisms (3x10e9) from a single rat (99-1-6,
sacrificed on 3/17/99) at Cincinnati VA facilities.
Trizol extracted RNA. Oligo dT priming, standard
conditions described by vendor, Stratagene. Further
details see www.uky.edu/Project/Pneumocystis/"
/organism="Pneumocystis
/mol_type="mRNA"
/db_xref="taxon:4754"
/lab_host="E. coli"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="mRNA"
/db_xref="taxon:4754"
/lab_host="E. coli"
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Pred. No. 54;
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                                                                                                                                                                                 of Kentucky, Lexington,
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COMMENT
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CF300133/c
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CF299342/c
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oryza sativa (japonica cultivar-group)
oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                 18;
CF300133 22 bp mRNA linear EST 15-AUG-2003 7LEAF--04-G19.bl Rice leaf plasmid cDNA library II (7LEAF) Oryza sativa (japonica cultivar-group) cDNA clone 7LEAF--04-G19, mRNA
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)

Contact: Nahm B.H.
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CF299342
CF299342.1 GI:33671103
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7LEAF--03-F06.b1 Rice leaf plasmid cDNA library II (7LEAF) Oryza
8ativa (japonica cultivar-group) cDNA clone 7LEAF--03-F06, mRNA
                                                                                                                                                                                                                                                                        Similarity
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: 82 31 330 6193
: 82 31 321 6355
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                   /tissue_type="leaf"
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/lab_host="E.coli DH10B"
/clone_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/clone_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
with oligoribonucleotides and then used as templates for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
/cultivar="Nackdong"
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'clone="7LEAF--03-F06"
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Pred. No. 54
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., L. Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., L.
Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Oryza sativa (japonica cultivar-group)
Cryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CF310366 22 bp mRNA linear EST 15-AUG.:
ABF--04-P14.gl ABF3-overexpressing transgenic rice plasmid cDNA
library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone
ABF--04-P14, mRNA sequence.
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Oryza sativa (japonica cultivar-group)
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                     Genomics and Genetics Institute, GreenGene Biotech Inc., of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                    Contact: Nahm B.H
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CF300133.1 GI:33671894
                                                                                      Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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                                                                   bhnahm@ggbio.com,
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/tissue_type="leaf"
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/note="Vector: pCR4-TOPO; Site_1: EcoR1; mRNA was capped
with oligoribonucleotides and then used as templates for
RT-PCR."
/organism="Oryza sativa
                                            Location/Qualifiers
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/mol_type="mRNA"
/cultivar="Nackdong"
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Pred. No. 54;
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                                                                   bhnahm@bio.myongji.ac.kr
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  (japonica cultivar-group)"
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CF311269/c
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nes 18; Conserv
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 22)
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22 bp mRNA linear EST 15-AUG-2003
ABF--06-G21.gl ABF3-overexpressing transgenic rice plasmid cDNA
library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone
ABF--06-G21, mRNA sequence.
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ilarity 81.8%;
Conservative
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Location/Qualifiers
                                                                                                                                                  /note="Vector: pCR4-TOPO; Site_1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Vector: pCR4-TOPO; Site_1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
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/dev stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone lib="ABF3-overexpressing transgenic rice plasmid
/cDNA library (ABF)"
                                                                                                                                                                                                                                                                            /dev_stage="14 days after germination"
/lab_host="5.coli DH108"
/clome lib="ABF3" overexpressing transgenic rice plasmid
cDNA library (ABF)"
                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Oryza sativa (japonica cultivar-group)"
|mol type="mRNA"
|culTivar="Nackdong"
|/db xref="taxon:39947"
|/clone="ABF--06-G21"
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/clone="ABF--04-P14"
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/cultivar="Nackdong"
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                            1.1%;
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Pred. No. 54;
0; Mismatches
Score 15.6; D
Pred. No. 54;
0; Mismatches
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                               Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 22)
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poacese;
Ehrhartoideae; Oryzaae; Oryza.
1 (bases 1 to 22)
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Lazge-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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CF311713.1 GI:33683474
Zin,
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Location/Qualifiers
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Similarity 81.8%;
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/dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone lib="ABF3-overexpressing transgenic rice plasmid cDNA library (ABF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; Leaf was dried for the color of the co
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/clone="ABF--07-B13"
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/cultivar="Nackdong"
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; Pred. No. 54;
0; Mismatches
Kim, M.J.,
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Lee, T.H., Shin, Y.C.,
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NACL--06-H22.bl Rice callus plasmid cDNA library (NACL) Oryza
sativa (japonica cultivar-group) cDNA clone NACL--06-H22, mRNA
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Contact: Nahm B.H.
                                                                                                                                                                                                                                            Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
Contact: Nahm B.H.
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopaida; Poalea; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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Location/Qualifiers
                                                                                                                                                                                                  bhnahm@ggbio.com, bhna
Location/Qualifiers
                                       /organism="Oryza sativa mol type="mRNA" (CULTivar="Nackdong" / db xref="taxon:39947" / Clone="NACL--06-H22" / tissue_type="callus" / tissue_type="callus"
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/lab_host="E.coli DH10B"
/clone_lib="ABF3-overexpressing transgenic rice plasmid
cDNA library (ABF)"
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/cultivar="Nackdong"
/db_xref="taxon:39947"
/clone="ABF--08-E15"
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dev_stage="proliferated callus"
| dev_host="E.coli DH10B"
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Pred. No. 54;
0; Mismatches
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CF334781 22 bp mRNA linear EST 18-AUG-20 JMT--04-D05.gl AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT) Oryza sativa (japonica cultivar-group) cDNA clone JMT--04-D05, mRNA sequence.
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                                                                                                                                                                                                                                                      18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of Bioscience and Bioinformatics, Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 22)
Xim,J.S., Jun,K.M., Cheong,P.J., Xim,M.J., Lee,T.H.,
Song,S.I., Xim,J.K., Xim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
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JMT--02-F04.gl AtJMT-overexpressing transgenic rice plasmid cDNA
library (JMT) Oryza sativa (japonica cultivar-group) cDNA clone
JMT--02-F04, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ehrhartoideae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bhnahm@ggbio.com, bhna
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                            /note="Vector: pCR4-TOPO; Site_1: EcoRI; Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was pepared from Arabidopsis Jasmoinate Carboxyl methyltransferase overexpression line."
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/note="Vector: pCR4-TOPO; Site 1: EcoRI; mRNA was capped
with oligoribonuclectides and then used as templates for
RT-PCR."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Oryza sativa
/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                      tissue type="leaf"
/dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone_lib="ALJMT-overexpressing transgenic rice plasmid
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/clone="JMT--02-F04"
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Pred. No. 54;
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Pred. No. 54;
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ACCESSION

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ORGANISM
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VERSION
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CF336250/c
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AUTHORS
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KEYWORDS
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JOURNAL
                                                                                                                                                                                                                                           TITLE
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Best Local Similarity
Matches 18; Conser
                                                                                                                                                                                                                                                                                          AUTHORS
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    source
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Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 22)
Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C.,
Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                 Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta;
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae;
Ehrhartoideae, Oryzeae, Oryza.
Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeongji, Korea Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
1. .22
                                                                                                                                                                                        1 (bases 1 to 22)

Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Kong, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)

Contact: Nahm B.H.
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22 bp mRNA linear EST 18-AUG-2003
JMT--66-D20.bl AtJMT-overexpressing transgenic rice plasmid cDNA
library (JMT) Oryza sativa (japonica cultivar-group) cDNA clone
JMT--06-D20, mRNA sequence.
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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EST.
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="leaf"
/dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone lib="AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT)"
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|mol_type="mRNA"
|/cultivar="Nackdong"
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clone="JMT--04-D05"
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Pred. No. 5
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1516 AATTAAAAAAAAAAAAGTAAAA 1537

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RESULT 77
CF337580
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  Matches
                    Query Match
Best Local
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Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbbo.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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                    1.1%; Score 15.6;
Similarity 81.8%; Pred. No. 5
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                                                                                                                                                                                                                                                                                      /mol type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:39947"
/clone="JMT--08-B11"
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/lab_host="5.coli DH10B"
/clone lib-"AtCMT-overexpressing transgenic rice
cDNA library (JMT)"
                                                                                                          CDNA library (JMT)"

/note="Vector: pCR4-TOPO; Site_1: EcoRI; Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was pepared from Arabidopsis Jasmoinate Carboxyl
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                                                                                       methyltransferase overexpression line."
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/clone_lib="AtJMT-overexpressing transgenic rice plasmid
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                                                                                                                                                                                                                                                               tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                                                              organism="Oryza sativa (japonica cultivar-group)"
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<u>.</u>
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                                         DB 1; Length 22;
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RESULT 79
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LOCUS
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                                    AUTHORS
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
1 (bases 1 to 22)
Abbal.P., Agasse,A., Ageorges,A., Atanassova,R., Barrieu,F.,
Couture,C., Dedaldechamp,F., Delrot,S., Glissant,D., Grimple
Hamdi,S., Romieu,C. and Terrier,N.
                                                                  Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledona; core eudicota; rosida; Vitaceae; Vitia.
                                                                                                                       Vitis vinifera
Vitis vinifera
                                                                                                                                                          CN545550.1 GI:46910175
EST.
                                                                                                                                                                                           EST 17494 Ripe Grape
clone B3CS00RL003D05
CN545550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oryza Bativa (japonica cultivar-group)
Oryza Bativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CF338524 22 bp mRNA linear E
RCL1--01-P07.g1 Regenerated callus lambda phage cDNA
Oryza sativa (japonica cultivar-group) cDNA clone RCI
                                                                                                                                                                                                                                                 CN545550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: bhnahm@ggbio.com, bhna
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yongin, Kyeonggi, K
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ehrhartoideae; Oryzeae; Oryza.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Vector: pBluescript SK(+); Site_1: Site_2: XhOI; CDNA was inserted into lamda Uni-ZAP XR Vector at end with SetI and 3' end with XhOI site. Callus was induced on 2N6 media for 30 days and cultured for 36hrs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               regenerated media"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
/cultivar="Nackdong"
/cultivar="Nackdong"
/db_xref="taxon:39947"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'Clone_lib="Regenerated callus lambda phage cDNA library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         dev stage="proliferated callus on 2N6 media for 30 days"
[ab_host="E.coli SOLR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              clone="RCL1--01-P07"
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                                                                                                                                                                                                                                                                                                                                                                                                                              81.8%;
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3', m
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Pred. No. 54
                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                            mRNA sequence.
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                  Grimplet, J.,
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VERSION
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AZ310066/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                         plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AZ310066.1 GI:10351682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seq primer: T7
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Similarity 81.88;
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                                                                                        Location/Qualifiers
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 db_xref="taxon:10090"
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Or seeus, ...
Unpublished (2002)
Contact: Hamdi S.
Contact: Hamdi S.
UMR 619 - Equipe Biologie de la Vigne
UMR 619 - Fordeaux I, Institut National
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                    Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ310066 22 bp DNA linear GSS 2: 1M0018D18R Mouse 10kb plasmid UUGCIM library Mus musculus clone UUGC1M0018D18 R, genomic survey sequence.
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                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0018 row: D column: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 00-33-(0)5-57-12-25-50
Fax: 00-33-(0)5-57-12-25-48
Email: s.hamdi@bordeaux.inra.fr
                                                                                                                                               Seq primer: CACACAGGAAACAGCTATGACCCClass: plasmid ends
                                                                                                                       quality sequence stop: 22.
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/clome_lib="Ripe Grape Skin Triplex2 Library"
/note="Organ: Fruit skin; Vector: Lambda TriplEx2; Site_1:
SfiIA; Site_2: SfiIB; Oriented library"
/mol_type="genomic DNA"
/strain="C57BL/6J"
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/culfivar="Cabernet Sauvignon"
/db xref="taxon:29760"
/clone="B3CS00RL003D05"
                                             organism="Mus musculus"
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                                                                                                                                                                                                                            Std Error: 0.00
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RESULT 81
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Best Local Similarity
Matches 18; Conserv
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0034 row: I column: 16
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Rese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       plasmid inserts
Unpublished (2000)
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                                                                                                                                              High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ316361.1 GI:10364110
                                                                                                                                                                                                       Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant valocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNPA2 (gil 4732114 [gb] AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and sheared adaptors to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone_lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42ny; Purified genomic DNA from M.
                        /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clone="UUGC1M0018D18"
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                                                                                 organism="Mus musculus"
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xref="taxon:10090"
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Pred. No. 54;
                                                                                                                                                                                                                                                                                         Std Error: 0.00
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FEATURES

COMMENT

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AUTHORS

KEYWORDS VERSION ACCESSION

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RESULT 82
AZ317017/c
LOCUS
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ORGANISM
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                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0035 row: P column: 09
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 22.
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Unpublished (2000)
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B. Weiss
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/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PMD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                /mol_type="genomic DNA'
/strain="C57BL/6J"
                                                                                                                                                    Location/Qualifiers
/db_xref="taxon:10090"
/clone="UUGC1M0035P09"
                                                                                              organism="Mus musculus"
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Pred. No. 5
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lab_host="E. Coli strain XL10-Gold, T1-resistant,

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JOURNAL COMMENT
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AUTHORS
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AZ351527
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VERSION
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Best Local (
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                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0089 row: E column: 07
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
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AZ351527
                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
Contact: Robert B. Weiss
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Location/Qualifiers
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/note="Vector: PWD42nv; Purified genomic DNA f
                                                                                                     /mol_type="genomic DNA"
/strain="CS7BL/6J"
                                              clone="UUGC1M0089E07"
                                                                               db_xref="taxon:10090"
                                                                                                                                                        organism="Mus musculus"
host="E. Coli strain XL10-Gold, T1-resistant,
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Insert Length: 10000 Std Error: 0
Plate: 0099 row: M column: 15
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 22.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Utah Genome Center 
University of Utah
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/note="Vector: PWD42nv; Purified genomic DNA f
                                                                                                                                                                                                                                                   /mol_type="genomic DNA"
/strain="C57BL/6J"
/lab_host="E. Coli strain XL10-Gold, T1-resistant,
/clone_lib="Mouse 10kb plasmid UUGCIM library"
                                                                                                                                                              clone="UUGC1M0099M15"
                                                                                                                                                                                                              /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                            organism="Mus musculus"
                                                                                                    sex="Male
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Pred. No. 54;
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ACCESSION VERSION

SOURCE KEYWORDS RESULT 84 AZ357630 LOCUS

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Matches

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AUTHORS
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AZ388103
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VERSION
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Best Local Similarity
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AZ388103
                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0147 row: N column: 14
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., 1slam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
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GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                         quality sequence stop: 22.
Location/Qualifiers
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/lab_host="B. Coli strain XL10-Gold, T1-resistant, F
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                               mol_type="genomic DNA"
strain="C57BL/6J"
                                                                                      sex="Male"
                                                                                                                                                                                                                          organism="Mus musculus"
                                                                                                               clone="UUGC1M0147N14"
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Pred. No. 5
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C57BL/6J (male) was obtained

REFERENCE AUTHORS

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RESULT 86
AZ401908/c
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ORGANISM
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Best Local (
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                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0168 row: P column: 24
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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1M0168P24R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University of Utah Genome Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert B. Weiss
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(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/lab_host="E. Coli strain XL10-Gold, T1-resistant,
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from N
                                                                                                                                                             /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                            ocation/Qualifiers
                                                                                     sex="Male"
                                                                                                            clone="UUGC1M0168P24"
                                                                                                                                            db_xref="taxon:10090"
                                                                                                                                                                                                                            organism="Mus musculus"
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Pred. No. 54;
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ACCESSION VERSION

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                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0203 row: A column: 24
Seq primer: CACACAGGAAACAGCTATGACC
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Mouse whole genome scaffolding with paired end reads from 10kb
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1M0203A24R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0203A24 R, genomic survey sequence.
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Contact: Robert B. Weiss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
/lab_host="B. Coli strain XL10-Gold, T1-resistant, /Clone_lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pW042nv; Purified genomic DNA from hmusculus C57BL/6J (male) was obtained from the Jac
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                                                                                                                                                                                        mol_type="genomic DNA"
/strain="C57BL/6J"
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'clone="UUGC1M0203A24"
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Pred. No. 54;
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Laboratory Mouse DNA Resource

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                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0212 row: A column: 05
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert B. University of Utah University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse_10kb_plasmid_UUGC1M_library"/note="Vector: pWd42nv; Purified_genomic_DNA_from M.musculus_CS7BL/6J (male) was obtained_from the Jackson_Laboratory_Mouse_DNA_Resource_(http://www.jax.org/resources/documents/dnares/). The DNA_from_the_dnares/).
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/strain="C57BL/6J"
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/clone="UUGC1M0212A05"
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                                                                                                                                                                           Bex="Male"
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Pred. No. 54;
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ACCESSION VERSION KEYWORDS

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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.

Mouse whole genome scaffolding with paired end reads from 10kb
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Inmert Length: 10000 Std Error: (
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Fax: 801 585 7177
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AZ459654.1 GI:10617779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                        ass: plasmid ends
gh quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse_10kb_plasmid_UUGCIM_library"
/note="Vector: PM042ny; Purified_genomic_DNA_from_M.
musculus_C57BL/6J_(male) was obtained_from_the_Jackson_Laboratory_Mouse_DNA_Resource
(http://www.jax.org/resources/documents/dnares/). The DNA_was hydrodynamically_sheared_by_repeated_passage_through_a
                                                                                                                                                                                                                                             /mol_type="genomic_DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                       clone="UUGC1M0264G12"
                                                                                                                                                                                                                                                                                                                           organism="Mus musculus"
                                                                                                                                                                                              sex="Male"
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Pred. No. 54;
0; Mismatches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0272 row: E column: 24
Seg primer: CGTTGTAAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ463503 22 bp DNA linear GSS 04-OCT-200 1M0272E24F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0272E24 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Class: plasmid ends
High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus (house mouse)
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/lab host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pWD42nv, Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                                                   /db_xref="taxon:10090"
/clone="UUGC1M0272E24"
                                                                                                                                                                                                                                                                                                                            mol_type="genomic DNA"
strain="C57BL/6J"
                                                                                                                                                                                                                                           sex="Male"
                                                                                                                                                                                                                                                                                                                                                                               organism="Mus musculus"
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31.8%; Pred. No. 54;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0272 row: E column: 12
Seq primer: CACACAGGAAACAGCTATGACC
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22 bp DNA linear GSS 0.
1M0272E12R Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC1M0272E12 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert B. Weiss
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/lab hosts "E. Coli strain XL10-Gold, T1-resistant, F-"
/clone libs "Mouse 10kb plasmid UUGCIM library"
/notes "Vector: PWD42nv; Furified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
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/clone="UUGC1M0272E12"
                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                             organism="Mus musculus"
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Pred. No. 54;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Nose whole genome scaffolding with paired end reads from 10kb
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Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0374 row: J column: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant vellocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were
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                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PW042TN', Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                              /db_xref="taxon:10090"
/clone="UUGC1M0374J15"
                                                                                                                                                                                                                                                                                                                                  /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                      /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                 organism="Mus musculus"
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Welss, R. Mouse, whole genome scaffolding with paired end reads from 10kb Mouse, whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0430 row: A column: 13
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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Fax: 801 585 7177
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University of Utah Genome Center
University of Utah
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/lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/clone="Vector: PMD42rv; Purified genomic DNA from M.
musculus C57BL/6y (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passes/ through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
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                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/strain="C57BL/6J"
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RESULT 94 AZ654691 VERSION KEYWORDS FEATURES REFERENCE ACCESSION DEFINITION TITLE JOURNAL ORGANISM Matches Query Match Best Local (1516 AATTAAAAAAAAAAAAGTAAAA 1537 18; 1 AAAAAAAAAAAAAAAAAAAA 22 Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Err Plate: 0529 row: D column: 0 Contact: Robert B. Weiss University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Research Bldg., 20 Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus. 1 (bases 1 to 22) AZ654691 22 bp DNA linear GSS 14 1M0529D05F Mouse 10kb plasmid UUGCIM library Mus musculus Clone UUGC1M0529D05 F, genomic survey sequence. Class: plasmid ends
High quality sequence stop: 22.
Location/Qualifiers Seq primer: CGTTGTAAAACGACGGCCAGT Unpublished (2000) Dunn, D., Aoyagi, A., Barber, M., AZ654691 Tel: 801 585 5606 Fax: 801 585 7177 plasmid inserts Mus musculus Mus musculus (house mouse) Similarity Conservative USA 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLL10-Gold (Stratagene) cells and selected for ampicillin resistance." (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGC1M library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource /organism="Mus musculus" /mol_type="genomic DNA" /strain="C57BL/6J" adaptored DNA was purified and size-selected for a 9.5 to /sex="Male" /db_xref="taxon:10090" clone="UUGC1M0529D05" 1.1%; Score 15.6; 81.8%; Pred. No. 54; 0; Mismatches Std Error: Beacorn, T., Duval, B., Hamil, C., 0.00 DB 1; 4. Length 22; <u>د</u> 2030 E., SLC, GSS 14-DEC-2000 0 Gaps genomic Ş 0

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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Nouse whole genome scaffolding with paired end reads from 10kb
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GSS.
Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0554 row: A column: 24
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligomucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel
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                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /Clone_lib="Mouse 10kb_plasmid_UUGCIM_Ibrary" /note="Vector: pWD42nv, purified_genomic_DNA_from_M. musculus C57BL/6J (male) was obtained_from_the_Jackson_musculus_C57BL/6J (mal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0554A24"
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; Pred. No. 54;
0; Mismatches
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2M0016I12R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0016I12 R, genomic survey sequence
AZ779844
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0016 row: I column: 1:
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University of Utah Genome Center
University of Utah
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 22)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: CACACAGGAAACAGCTATGACC
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801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative
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                                                                                                                                                                                                                  Biomedical Polymers Research Bldg., 20 S. 2030
                                                                                                                                                                                                                                                                                                                                                                                                                /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: C
Plate: 0028 row: E column: 04
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 22)
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Mus musculus (house mou
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2M0028E04R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0028E04 R, genomic survey sequence.
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Fax: 801 585 7177
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gi|4732114|gb|AF129072.1), a copy-number
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                                                                                                                                                                                                                                                                                        /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                               clone="UUGC2M0028E04"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="Mus musculus"
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Pred. No. 54;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.eu
Insert Length: 10000 Std E
Plate: 0033 row: A column:
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
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Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polymucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gii 4732114 [gb|AF119072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
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/clone="UUGC2M0033A05"
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/strain="C57BL/6J"
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81.8%; Pred. No. 54;
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Insert Length: 10000 Std Error:
Plate: 0034 row: G column: 12
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University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes R., Tingey, A., von Niederhausern, A. and Wright, D., Weise, R.
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High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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GSS.
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Conservative
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNB42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
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                                                                                                                                                                                                                                                                                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             clone="UUGC2M0034G12"
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Pred. No. 54;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0045 row: A column: 2
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University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Class: plasmid ends
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Fax: 801 585 7177
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http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNA92 (gi|473114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="genomic
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                              sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       clone="UUGC2M0045A24"
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Pred. No. 54;
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Insert Length: 10000 Std Error:
Plate: 0076 row: E column: 19
Seg primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
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ilarity 81.8%;
Conservative
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             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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/clone_lib="Mouse_10kb_plasmid_UUGCIM_library"
/note="Vector: pM042nv, Purified_genomic_DNA_from M.
musculus_C578L/6J_(male) was obtained_from the Jackson
Laboratory_Mouse_DNA_Resource_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biomedical Polymers Research Bldg.,
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adaptored vector DNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     clone="UUGC2M0076E19"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                db_xref="taxon:10090"
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 Mismatches

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  transformed into
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Query Match
Best Local (
                                                       chemically-competent E. coli XL10-Gold and selected for ampicillin resistance.
               1.1%;
Score 15.6; DI
Pred. No. 54;
O; Mismatches
                              DB 1;
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Matches

18; Conservative

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AZ820439/c
LOCUS
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KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Class: plasmid ends
High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 092 row: K column: 13
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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2M0092K13R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0092K13 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
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/lab host="E. Coli strain Xilo-Gold, T1-resistant, F-"
/clone_lib="Mouse lokb plasmid UUGCIM library"
/note="Vector: PMD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polymucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
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/clone="UUGC2M0092K13"
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sex="Male"
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AZ841661
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Best Local
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0139 row: I column: 18
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Utah Genome Center University of Utah
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GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B. Weiss
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGC1M library"/note="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackso
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clone="UUGC2M0139I18"
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AUTHORS
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 22)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Nouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0142 row: I column: 24
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2M0142I24F Mouse 10kb plasmid UUGC1M library Muclone UUGC2M0142I24 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ843514.1 GI:13013422
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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pM042 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/Clome_lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PW927v; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone="UUGC2M0142I24"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="Mus musculus"
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Pred. No. 54;
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AZ946102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rockes, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0207 row: D column: 13
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                    adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid RI. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored.
adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                           (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant vellocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                             /lab host="E. coli strain XL10-Gold, T1-resistant, F-"
/clome_lib="mouse 10kb plasmid UUGC2M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C7BL/6J (female) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clone="UUGC2M0207D13"
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sex="Female"
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Pred. No. 5;
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Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18; Conservative
                                                                              Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           insert libraries for whole genome shotgun sequencing projects. In Genome Sequencing: A practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).

Email: nelsayed@tigr.org
Details nelsayed@tigr.org at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.

Location/Qualifiers
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Constructed at the Institute for Genomic Research (TIGR),
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shear
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shear
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described in detail in Smith, H. and Venter, J.C. (Making small
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Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L.,
                                                                                                                                                        GSS.
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/strain="TREU927"
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A clone 131b09, forward sequence,
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                                                                 Bource
                                                                                                                                                     Direct Submission

L Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CBIO 1SA, E-mail: barrell@sanger.ac.uk and nhl@sanger.ac.uk

Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared to give a tight size distribution (

4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
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4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
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                                                                                Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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Location/Qualifiers
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/mol_type="genomic DNA"
/strain="TREU927"
/organism="Trypanosoma brucei"
/mol_type="genomic DNA"
/strain="TREU927"
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DNA clone 35c12, reverse sequence
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AG194579.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 ISA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Trypanosoma brucei
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                     Pan troglodytes (chimpanzee)
Pan troglodytes
                                                                                                                                        sequence.
                                                                                                                                                                                     AG194579
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Similarity 81.8%;
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/clone="35c12"
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/clone="380a07"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mol_type="genomic DNA"
/strain="TREU927"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Trypanosoma brucei"
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Pred. No. 54;
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CF334610/c
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Best Local Similarity
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                                                                                                                                                                                                           Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzaae; Oryza.

1 (bases 1 to 19)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                          CF334610
CF334610.1 GI:33817556
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CF334610 19 bp mRNA linear EST 18-AUG-2003 JMT--03-P13.bl AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT) Oryza sativa (japonica cultivar-group) cDNA clone JMT--03-P13, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC); 52, Oun-dong, Yusong-gu, Daejeon 305-333, Korea (E-mail:redstone@mail.kribb.re.kr, URL:http://phs.grc.kribb.re.kr/, Tel:82-42-866-7181, Fax:82-42-860-4409)
Clones are derived from the chimpanzee BAC library RP-43 This BAC end was generated during the R&D process and may have higher chance
                                                       Email: bhnahm@ggbio.com, bhn.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of clone tracking errors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Park, K., Eun, C.J., Hoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C., Song, W. and Yoo, H. BAC end sequences of Library RP-43
Unpublished
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                                                                                               Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                       Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                               Contact: Nahm B.H
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R.Site 1 : EcoRI
R.Site 2 : EcoRI.
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/clone_lib="RP-43 Chimpanzee Male BAC Library"
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/mol_type="genomic DNA"

/db_xref="taxon:9598"
/organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone="RP43-072N05.TJ"
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                                                                                                                                       Korea
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Pred. No. 54;
                                                                               bhnahm@bio.myongji.ac.
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       CF319122
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CL680297
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Max-Planck-Institute for Developmental Biology
Spemannstr. 37-39, Tuebingen D-72076, Germany
Tel: 00497071601371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Srinivasan,J., Otto,G.W., Kahlow,U., Geisle
AppaDB: an AcedB database for the nematode
Pristionchus pacificus
Nucleic Acids Res. 32 (1), D421-D422 (2004)
Contact: Sommer RJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pristionchus pacificus
Eukaryota; Metazoa; Nematoda; Chromadorea; Diplogasterida;
Neodiplogasteridae; Pristionchus.
1 (bases 1 to 20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ralf.sommer@tuebingen.mpg.de
This library was generated at Caltech,
sequenced at Vancouver, Canada.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Class: fosmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fax: 00497071601498
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cDNA_library (JMT)"

/note="Vector: pcr4-TOPO; Site_1: EcoRI; Oligo-capped mRNA
/note="Vector: pcr4-ToPO; Oligo-capped mRNA
/note="Vector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Pristionchus
/mol_type="genomic DNA"
/strain="California"
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/lab_host="E.coli DH10B"
/clone_lib="AtJMT-overexpressing transgenic rice plasmid
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/db_xref="taxon:39947"
/clone="JMT--03-P13"
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var. California"
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94.1%; Pred. No. 53;
tive 0; Mismatches
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Pred. No. 56;
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for the nematode satellite
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15-AUG-2003

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                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 25)
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749097
755 bp mRNA linear EST 06-FEB-19
7b08h08.81 Stratagene placenta (#937225) Homo sapiens cDNA clone
IMAGE.70623 3' similar to similar to gb.%62744 CLASS II
HISTOCOMPATIBILITY ANTIGEN, M ALPHA CHAIN (HUMAN), mRNA sequence.
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae,
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CF319122
Generation and analysis of 280,000 human Genome Res. 6 (9), 807-828 (1996)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Large-scale Sequencing Analysis 
Unpublished (2003)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="callus"
/dev_stage="proliferated callus on 2N6 media for 2 weeks"
/lab_host="E.coli DH10B"
/clome_lib="OsHDAC1-overexpressing transgenic rice plasmid cDNA library (HD)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Vector: pCR4-TOPO; Site_1: EcoRI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was
                                                                                                                                                                                                                                                                                                                                          GI:650957
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/mol_type="mRNA"
/cultivar="Nackdong"
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clone="HD--09-107"
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O; Mismatches
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                    expressed sequence tags
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Homo sapiens
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High qality sequence stops: 1
Source: IMAGE Consortium, LINL
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
                                                                                                                                                                                           Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany
    Location/Qualifiers
                                                                                                                                                                                                                                                    Contact: MIPS
                                                                                                                                                                                                                                                                     EST (Ottenwaelder, 
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                          Ottenwaelder, B., Obermaier, B., Mewes, H.W.,
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High quality sequence stop: 1.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Other_ESTs: yb08h08.rl
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                           Wiemann, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                               AL038460.1 GI:49682131
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314 286 1810
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/organism="Homo sapiens"
/mol type="mRNA"
/db xref="taxon:9606"
/clone="pkrzp56682246"
/tissue_type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMPl; Site_1: NotI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /lab_host="SOLR cells (kanamycin resistant)"
/clone_lib="Stratagene placenta (#37225)"
/note="Organ: placenta; Vector: pbluescript SK-; Site_1:
EcoRI; Site_2: XhoI; Cloned unidirectionally. Primer:
Oligo dT. Caucasian. Average insert size: 1.2 kb; Uni-ZAP
XR Vector; -5' addaptor sequence: 5' GAATTCGGCACGAG 3' ~3'
adaptor sequence: 5' CTCGAGTTTTTTTTTTTTTTTT 3'"
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/mol_type="mRNA"
/db_xref="gDB:491520"
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/clone="IMAGE:70623"
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Pred. No. 69;
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onym: hfkd2) Homo
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       NotI; Site_2: SalI"
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RESULT 117
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Oryza sativa (japonica cultivar-group)
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
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                                                          CF339443
CF339443.1 GI:33827271
EST.
                                                                                                               CF339443 20 bp mRNA linear EST 18-AUG-2003 RCL1--04-003.gl Regenerated callus lambda phage cDNA library (RCL1) Cryza sativa (japonica cultivar-group) cDNA clone RCL1--04-003,
                                                                                                                                                                                                                                                                              l Similarity
17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 20)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7LEAF--01-D19.b1 sativa (japonica
                                                                                                    mRNA sequence.
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                                                                                                                                                                                                                                               <u> АДТТАДАДАДАДАДАДСТДА</u> 1535
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nilarity 85.0%;
Conservative
                                                                                                                                                                                                                                                                                                                              Conservative
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="Oryza sativa (japonica cultivar-group)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GI:33669779
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Pred. No. 63;
O; Mismatches
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Pred. No. 63;
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AJ661013/c
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AJ661013
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                        Contact: Anderson SI
Genomics and Bioinformatics
                                                                                                                                                                                                                                                                                                                                        Sus scrofa
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/cultivar="Nackdong"
/db_xref="taxon:39947"
                               organism="Sus scrofa"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tissue_type="callus"
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Pred. No. 63;
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Roslin Institute
Roslin, Midlothian, EH25 9PS, UNITED KINGDOM
Roslin, Midlothian, EH25 9PS, UNITED KINGDOM
Single pass sequencing. Bases called and trimmed with phred
VO.020425 c. Vector identified by cross match with the -minscore 20
and -minmatch 12 options. Vector:pBlueScriptII(KS+) R. Site 1:
ECORI R. Site 2: NotI Description: Normalised library constructed
from pooled tissue from day 30 placentas. Clones available from UK
Centre for Functional Genomics in Farm Animals, Roslin Institite,
Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae;
1 (bases 1 to 21)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anderson, S.I., Finlayson, H.A. and Archibald, A.L. Development of cDNA and EST resources for studying reproduction and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AJ661013 CSEQRAN09 Sus scrofa cDNA clone C0000935_H04, mRNA
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|lab_host="E_coli SOLR"
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(RCL1)<sup>m</sup>
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Sus.
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AL038582
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Best Local
                                 AUTHORS
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Wiemann,S.
EST (Otten
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DKFZp566H2046_r1 566 (synonym:
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1 (bases 1 to 21)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AL038582 21 bp mRNA linear EST 06-JUI DKFZp566F0946 rl 566 (synonym: hfkd2) Homo sapiens cDNA clone DKFZp566F0946, mRNA sequence.
AL038582
                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 21)
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Location/Qualifiers
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Unpublished (1999)
                             Ottenwaelder, B., Obermaier, B.,
                                                                                                    Homo sapiens
                                                                                                                                                  AL038627.1 GI:49682173
                                                                                                                                                                                 DKFZp566H2046, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: MIPS
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(Ottenwaelder,
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/db_xref="taxon:9606"
/clone="bkr2p566F0946"
/tissue_type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: Not1; Site_2: Sa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="placenta"
/clone_Tib="CSEQRANO9"
/note="Vector: pBlueScriptII(KS+); Site_1: EcoRI; Site_2:
NotI; Single pass sequencing. Normalised library
constructed from pooled tissue from day 30 placentas."
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~nvm: hfkd2) Homo
                                 Mewes, H.W., Gassenhuber, J. and
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Best Local
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sequence.
CF330439
                        CF330439 21 bp mRNA linear EST 18-AUG-2003
NACL--06-C12.b1 Rice callus plasmid cDNA library (NACL) Oryza
sativa (japonica cultivar-group) cDNA clone NACL--06-C12, mRNA
                                                                                                                                 1 TCARAAAAAAAAAAAAAAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17;
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Contact: MIPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DKFZp566P1346 r1 566 (synonym: hfkd2) Homo sapiens DKFZp566P1346, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                  Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: MIPS
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Unpublished (1999)
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Mammalia; Eutheria;
1 (bases 1 to 21)
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="DKFZp566H2046"
/tissue_type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: Not1; Si
                                                                                                                                                                                                                                              /mol_type="mRNA"
/db_xref="taxon:9606"
/clone="bKrZp56661346"
/tissue_type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI;
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Primates;
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Pred. No. 66;
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CDNA clone
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Eukaryota; Viridiplantae; Streptophyta; Embryor
Spermatophyta; Magnoliophyta; Liliopsida; Poale
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 21)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17;
                               UMR BiO3P, BP 35327, F-35653 Le Rheu Cedex France
Tel: +33.2.23.48.51.65
Fax: +33.2.23.48.51.50
Risk of contamination by bacterial sequences from obligatory
(Buchnera) or facultative endosymbionts. These sequences were
obtained in the frame of the International Consortium of Aphid
Genomics in collaboration with Genoscope
                                                                                                                                                                                                 Unpublished (2004)
Contact: D. Tagu
                                                                                                                                                                                                                                                          Hunter, W., Martinez-Torres, D., Rahbe, Y., Sabater-Munoz, B., Stern, D., Tagu, D. and Wincker, P.
An expressed sequence tags database for the pea aphid Acyrthosiphon
                                                                                                                                                                                                                                                                                                                     Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota, Neoptera, Paraneoptera, Hemiptera, Sternorrhyncha, Aphidiformes, Aphidoidea, Aphididae, Macrosiphini, Acyrthosiphon.

[ (bases 1 to 21)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mRNA sequence.
CN763587
CN763587.1 GI:47537510
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EST.
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Acyrthosiphon pisum
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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FORWARD: CAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:39947"
/clone="NACL-06-C12"
/tissue_type="callus"
/dev_stage="proliferated callus on 2N6 media for 30 days"
/lab_host="E.coli DH10B"
/clone lib="Rice callus plasmid cDNA library (NACL)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped with oligoribonucleotides and then used as templates for RT-PCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
/cultivar="Nackdong"
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a; Poales; Poaceae;
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                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0156 row: F column: 13
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 21)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ393269 21 bp DNA linear GSS 03-OCT-200100156F13F Mouse 10kb plasmid UUGC1M library Mus musculus genomic Clone UUGC1M0156F13 F, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                             University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plasmid inserts
Unpublished (2000)
                                                                                                                                                            Class: plasmid ends
High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B. Weiss
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                                                                                                                                                                                                                                                                                                                                                                          Biomedical Polymers Research Bldg.,
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Location/Qualifiers
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                                         /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                 /organism="Mus musculus"
/mol_type="genomic DNA"
                                                                                                                                         Location/Qualifiers
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/note="Vector: pB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /dev_stage="nymphs and adults (parthenogenetic females)"
/lab_host="XL1-Blue"
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/cultivar="developmentstage"
                        /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tissue_type="whole insect"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      clone="IDOAAA7BH12"
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clone="UUGC1M0156F13"
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Pred. No. 66
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lab_host="E. Coli strain XL10-Gold, T1-resistant,

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ORGANISM
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KEYWORDS
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AZ625662/c
                                                                                                                                                                                                      FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0465 row: C column: 23
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota, Metazoa; Chordata; Craniata; Vercebrata; Euceleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 21)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 AATTAGTATTTTTTAATTTT 20
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GSS.
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                                                                                                                                                                                                                        High quality sequence stop: 21.
                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical Polymers Research Bldg.,
                                                                                             organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                 ocation/Qualifiers
                            'sex="Male"
                                                 clone="UUGC1M0465C23"
                                                                              db xref="taxon:10090"
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host-"E. Coli strain XL10-Gold, T1-resistant,
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Pred. No. 66;
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DEFINITION
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CF300172
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Ewkaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CF300172 23 bp mRNA linear EST 15-AUG-2003 7LEAF--04-H15.b1 Rice leaf plasmid cDNA library II (7LEAF) Oryza sativa (japonica cultivar-group) cDNA clone 7LEAF--04-H15, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Nahm B.H.

Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                      Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/60 (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi 4732114|gb|AFT29072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                               bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
/dev_stage="7 days after germination"
/lab_host="E.coli DH10B"
/clone_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/clone_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was cappo
with oligoribonucleotides and then used as templates for
                                                                                                                                             /organism="Oryza sativa (japonica cultivar-group)"

mol type="mRNA"

/culTivar="Nackdong"

/db xref="taxon:39947"

/clone="7LEAF--04-H15"

/tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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Pred. No. 66;
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Query Match Best Local Similarity

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                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AL Unpublished (1999)
Other EST8: 2822740.5prime
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Email: cgapbs-r@mail.nih.gov
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
Tissue Procurement: DCTD/DTP cDNA Library Preparation: Ling
Consortium (LLNI) DNA Sequencing by Berkeley MGC sequencing
project Clone distribution: MGC clone distribution information can
be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrg/image/inage.html Base Calling / Quality
Scores: PHRED from University of Washingtion Genome Center. Vector
Trimming: cross match from University of Washingtion Genome Center
PHRAP suite- Poly-T Identification: patMatch.pl from Berkeley
Drosophila Genome Project: University of Washingtion Genome Center:
http://www.genome.washington.edu Low Quality Sequence: Genter:
http://www.genome.washington.edu Low Quality Sequence: Trace file contained 15 contiguous distinct peaks
following vector sequence. Polyadenylation: Based upon the presence
of a XhoI site followed by a run of 14 or more T residues at the
beginning of the sequence, this cDNA insert was polyadenylated.
Plate: LLCM10 row: D column: 5
High quality sequence stop: 6.
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1 (bases 1 to 15)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
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2822740.3prime NIH_MGC_7 Homo sapiens cDNA clone IMAGE:2822740 3',
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/clone lib="NIH MGC_7"
/note="Organ: lung; Vector: pOTB7; Site 1: XhoI; Site 2: EcoRI; CDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Limg Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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/tissue_type="small cell carcinoma"
/cell_line="MGC3"
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100.0%; Pred. No.
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Insert Length: 10000 Std Error:
Plate: 0004 row: P column: 01
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ774536 19 bp DNA linear GSS 10 2M0004P01F Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0004P01 F, genomic survey sequence.
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Location/Qualifiers
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University of Utah Genome Center
University of Utah
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Fax: 801 585 7177
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                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                          /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/GJ (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pMs042 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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ACCESSION: AR093142 ACCESSION: CAB30491 ACCESSION: AR279103 ACCESSION: AX183931 ACCESSION: AX183931 ACCESSION: AX1825110 ACCESSION: AX825114 ACCESSION: AX825118 ACCESSION: AX825112 ACCESSION: AX825122 ACCESSION: AX825138 ACCESSION: AX825134 ACCESSION: AX825134 ACCESSION: AX825154	ACCESSION: A01996 ACCESSION: A0442 ACCESSION: AR294636 ACCESSION: AR084566 ACCESSION: AR084579 ACCESSION: AR084579 ACCESSION: AR084579 ACCESSION: AR084579	ACCESSION: AR431307 ACCESSION: AR253000 ACCESSION: AR253103 ACCESSION: AX825104 ACCESSION: AX825105 ACCESSION: AX825115 ACCESSION: AX825151	ACCESSION: AX042782 ACCESSION: AR370697 ACCESSION: AX117632 ACCESSION: AX692829 ACCESSION: AX692830 ACCESSION: AX692831 ACCESSION: AX692831 ACCESSION: AX692831	Description	y chance to have a e result being printed, distribution.			964		cataactaatctcttctgtt 1401	me 7 Seconds gnments) n cell updates/sec	Ltd.
												
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SOURCE Unknown. SOURCE Unknown. ORGANISM Unknown. Unclassified. VINCLASSIFIED. REFERENCE 1 (bases 1 to 25) AUTHORS Barbas, C.F., Burton, D.R. and Lerner, R.A. AUTHORS Synthetic human neutralizing monoclonal antibodies to human immunodeficiency virus JOURNAL Patent: US 6395275-A 10 28-MAY-2002; FEATURES Location/Qualifiers Source 125	Db 1 TritritritritaddAcAcAdArCTT 25 RESULT 2 AR370697/c AR370697	/ Operation	synthetic construct synthetic construct artificial sequences. 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	ALIGNMENTS RESULT 1 AX042782 LOCUS AX042782 DEFINITION Sequence 348 from Patent WO0065088. ACCESSION AX042782 VERSION AX042782 VERSION AX042782.1 GI:11341390	472 14.2 1.0 20 1 AR489489 473 14.2 1.0 20 1 AR791100 ACCESSION: AR4891100 C 475 14.2 1.0 20 1 AX078001 C 476 14.2 1.0 20 1 AX078001 C 477 14.2 1.0 20 1 AX37428 C 477 14.2 1.0 20 1 AX293619 C 478 14.2 1.0 20 1 AX293619 C 479 14.2 1.0 20 1 AX293619 C 480 14.2 1.0 20 1 AX452909 C 480 14.2 1.0 20 1 AX452909 C 481 14.2 1.0 20 1 AX452909 C 482 14.2 1.0 20 1 AX452909 C 483 14.2 1.0 20 1 AX750458 C 483 14.2 1.0 20 1 AX785487 C 485 14.2 1.0 20 1 AX785487 C 486 14.2 1.0 20 1 AX785487 C 487 14.2 1.0 20 1 AX785487 C 488 14.2 1.0 20 1 BD056369 C 488 14.2 1.0 20 1 BD056369 C 489 14.2 1.0 20 1 BD056369 C 499 14.2 1.0 20 1 BD056309 C 499 14.2 1.0 20 1 BD056369 C 499 14.2 1.0 20 1 BD056309 C 490 14.2 1.0 20 1 BD056309

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                                                                                                                                                                                                    Shannon, M., Gu, Y. and Nguyen, C.T. Four human zinc-finger-containing proteins: mdz3, mdz4, mdz7
                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                /organism="Homo sapiens"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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RESULT 7
AX825106/c
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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                                                                                                                                                                                            1 (bases 1 to 24)
Vaisberg, E.A., Adams, C.L., Sabry, J.H. and Crompton, A.M.
Database system including computer code for predictive cellular
                                                                                                                                                                                                                                                                                                            Sequence 1 from patent US AR431307
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                                                                                                          /organism="unknown"
/mol_type="genomic DNA"
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/mod_base=OTHER
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/mod_base=OTHER
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Penny,L. and Galvin,M.
Genotyping the human UDP-glucuronosyltransferase 1
Patent: US 6479236-A 100 12-NOV-2002;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                              /organism="synthetic construct"
/mol_type="unassigned DNA"
/db xrefs"ttaxon:32630"
/noTe="Beschreibung der kuenstlichen
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                                                             /mod_base=OTHER
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                                                                                                   mod_base=OTHER
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Sequence 2 from Patent WO03072818.
AX825104
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Sequence 3 from Patent WO03072818.
AX825105
AX825105.1 GI:39750834
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Boekenkamp,D., Dieck,T.H. and Hoppe,H. Method for sorting single-stranded nuc
                                        synthetic construct synthetic construct artificial sequences.
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Sequenz:Capture-Oligonukleotid"
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Sequence 49 from Patent WO03072818.
AX825151
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Degussa Bioactives GmbH (DE)
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llarity 90.0%; Pred. No. 53;
Conservative 0; Mismatches
  /mod_base=OTHER
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/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
                                                      /note="LNA-T (Locked
/mod_base=OTHER
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mod_base=OTHER
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mol_type="unassigned DNA"
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Chetverin,A.B. and Kramer,F.R.
Oligonucleotide arrays and their use for sorting,
sequencing, and manipulating nucleic acids
patent: US 6322971-A 6 27-NOV-2001;
Location/Qualifiers
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                                                                                                                                     Patent: WO 8404538-A 24 22-NOV-1984;
Location/Qualifiers
                                                                                                                                                                     synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 23)
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/mod_base=OTHER
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                                                                                        /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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                                           Score 16.6; Di
Pred. No. 55;
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Pred. No. 46;
0; Mismatches
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Cohen,D., Chumakov,I. and Blumenfeld,M.
Biallelic markers for use in constructing
disequilibrium map of the human genome
Patent: US 6537751-A 6371 25-MAR-2003;
Location/Qualifiers
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1 (bases 1 to 23)
Edens,L., Russell,S.W., Visser,C.
Improvements in the expression of cells
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Unclassified.
1 (bases 1 to 21)
                       Unknown.
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UNILEVER NV; UNILEVER PLC
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Similarity 94.4%; Pred. No. 75;
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6371 from patent US
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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Sequence 67
AR084578
                                           18;
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                                                                                                                 1 (bases 1 to 21)
Matson,R.S., Coassin,P.J., Rampal,J.B. and
Oligonucleotide repeat arrays
Patent: US 5981185-A 68 09-NOV-1999;
Location/Qualifiers
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Matson,R.S., Coassin,P.J., Rampal,J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 67 09-NOV-1999;
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/mol_type="unassigned DNA"
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Pred. No. 76;
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AR084582
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Sequence 3 from Patent WO2004055153.
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Matson, R.S., Coassin, P.J., Rampal, J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 71 09-NOV-1999;
Location/Qualifiers
Schluesener,H. and Wendel,H.P. Devices coated with substances that mediate biological material Patent: WO 2004055153-A 3 01-JUL-2004; Eberhard-Karls-Universitaet Tuebingen (DE)
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Bergan,R. and Neckers,L.
Inhibition of protein kinase activity by aptameric action of
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AR093142
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                                                                    artificial sequences.
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/mol_type="unassigned
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Sequence 34 from Patent W00214546.
AX383931
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Patent: WO 0214546-A 34 21-FEB-2002;
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Plasmodium falciparum
Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
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1 (bases 1 to 21)
Milhausen, M.J.
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                                                                                                   /organism="Plasmodium
/mol_type="unassigned
/db_xref="taxon:5833"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="unknown"
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db_xref="taxon:32630"
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AX713257/c
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                                                                                                                                                                                                                                                                                                                                                                           synthetic construct synthetic construct artificial sequences.
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/mol type="unassigned DNA"
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/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/note="LNA-T (Locked
/mod_base=OTHER
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                                                     note="LNA-T (Locked
                                                                                                                                note="LNA-T (Locked Nucleic Acid)"
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/mol type="unassigned DNA"
/db_xref="texon:32630"
/note="Beschreibung der kuenstlichen
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Location/Qualifiers
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Sequence 20 from Patent WO03072818
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/mol type="unassigned DNA"
/mbxref="taxon:32630"
/nbte="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
 /note="LNA-T (Locked Nucleic Acid)"
                                                                                                  /note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
                                                                                                                              organism="synthetic construct"
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/mod_base=OTHER
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/mod_base=OTHER
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Sequence 36 from Patent W003072818.
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mod_base=OTHER
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JP 2001333800-A/1.
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                                     Method of comparison and detection of RNA Patent: JP 2001333800-A 1 04-DEC-2001; UNITECH CO LTD
                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                   Homo sapiens (human)
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Homo sapiens (human)
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/mb xref="taxon:32630"
/note="Beschreibung der kuenstlichen
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Sequence
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                                                                    1 (bases 1 to 20)
Vasseur,M., Blumenfeld,M., Meguenni,S. and Poddevin,B.
STAPLE AND SEMI-STAPLE OLIGONUCLEOTIDES, METHOD OF PRE
APPLICATIONS
Patent: WO 9423026-A 2 13-OCT-1994;
GENSET (FR)
                                                                                                                                                                                                         20 bp
Sequence 2 from Patent WO9423026.
A40126
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1 (bases 1 to 20)
Griffais,R., Hoiseth,S.K., Zagursky,R.J., Metcalf,B.J., Peek,J.A., Sankaran,B. and Fletcher,L.D.
Chlamydia pneumoniae polynucleotides and uses thereof Patent: US 6559294-A 6233 06-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N
                                                                                                                                              unidentified unclassified.
                               Other publication AU 6432094 941024
Other publication FR 2703053 940930
Location/Qualifiers
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C12Q1/68,C12N15/09,G01N33/50,C12N15/00
Method of comparison and detection of RNA amount and
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/organism="unidentified"
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/mol_type="genomic RNA"
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Location/Qualifiers
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/mol_type="genomic DNA"
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Pred. No. 73;
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AR139961/c
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AR140280/c
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AR140558
Sequence 33 from
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Zsebo,K.M., Bosselman,R.A., Suggs,S.V. and Martin,F.H.

Method for enhancing the effciency of gene transfer with stem cell
factor (SCF) polypeptide
Patent: US 6207454-A 33 27-MAR-2001;
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Zsebo, K.M., Bosselman, R.A., Suggs, S.V.
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/mol_type="unassigned DNA"
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/mol_type="unassigned
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/db_xref="taxon:32644"
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Pred. No. 1e+02;
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Pred. No. 1e+02;
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AR182885
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1 (bases 1) to 20)
Krieg,A.M., Davis,H.L., Wu,T. and Schorr,J.
Vectors and methods for immunization or therapeutic protocols
Patent: US 6339068-A 57 15-JAN-2002;
Location/Qualifiers
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Zsebo, K.M., Bosselman, R.A., Suggs, S.V.
Stem cell factor and compositions
Patent: US 6207802-A 33 27-MAR-2001;
                                                                       Krieg, A.M., Schetter C. and Vollmer, J.C. Immunostimulatory nucleic acids Patent: WO 0122972 nucleic Scher-2001; UNIVERSITY OF IOWA RESEARCH FOUNDATION (US); Coley Pharmaceutical
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synthetic construct
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organism="synthetic construct"
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/db_xref="taxon:32630"
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Sequence
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UNIVERSITY OF IOWA RESEARCH FOUNDATION
Location/Qualifiers
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Similarity 89.5%;
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GGCGGCGGCGGCGGCG
                 GGCGGCTGCGGCGCCGGCG
                                             1.1%;
larity 89.5%;
Conservative
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                                                                                                                                                                                                        for enhancing antibody-induced cell
                                                                                                                                                                                                                                                                                                                    410 from Patent
                                                                                             /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="caxon:32630"
/noTe="Synthetic oligonucleotide~phosphodiester
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="synthetic construct"
/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noTe="Synthetically generated primer"
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                      444
                                             Score 15.8; DB 1;
Pred. No. 1e+02;
0; Mismatches 2;
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19
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Pred. No. 1e+02;
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BD069976
LOCUS
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VERSION
KEYWORDS
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ACCESSION
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Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                              Schwartz,D.A. and Krieg,A.M.

Use of nucleic acids containing unmethylated CPG dinucleotide in the treatment of LPS-associated disorders

LP 2001513776-A 65 04-SEP-2001;

UNIVERSITY OF 10WA RESEARCH FOUNDATION

OS Artificial Sequence
PN JP 2001513776-AA65
PD 04-SEP-2001
PF 25-PEB-1998 JP 1998537810
PF 26-PEB-1999 UP 1998537810
PF 26-PEB-1999 UP 1998537810
PF 26-PEB-1999 UP 1998537810
PF 26-PEB-1999 UP 1998537810
PF 26-PEB-1990 UP 1998537810
PF 26-
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Bratzler,R.L.

Inhibition of angiogenesis by nucleic acids Patent: WO 02053141-A 243 11-UL-2002; Coley Pharmaceutical Group, Inc. (US)

Location/Qualifiers
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1 (bases 1 to 20)
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JP 2001513776-A/65.
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Use of nucleic acids containing the treatment of LPS-associated
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GGCGGCGGCGGCGGCG 19
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                                                                                                                                                                                                                                                                 /organism='Artificial Sequence'
Location/Qualifiers
                                                                                                                                                                                  /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Pred. No. 1e+0
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AR084567
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AUTHORS
TITLE
JOURNAL
FEATURES
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CQ830490/c
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DEFINITION
ACCESSION
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FEATURES
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AR084563/c
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Sequence 56
AR084567
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Sequence 52 from patent US 5981185.
AR084563
AR084563.1 GI:10011374
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Sequence 2 from Patent WO2004055153.
C0830490
CQB30490.1 GI:50250830
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1 (bases 1 to 21)

Natson,R.S., Coassin,P.J., Rampal,J.B. and Caskey,C.Thomas.
Oligonucleotide repeat arrays
Patent: US 5981185-A 52 09-NOV-1999;
Location/Qualifiers
Schluesener,H. and Wendel,H.P. Devices coated with substances that mediate biological material Patent: WO 2004055153-A 2 01-JUL-2004; Eberhard-Karls-Universitaet Tuebingen (DE)
                                                                              synthetic construct artificial sequences.
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Matson,R.S., Coassin,P.J., Rampal,J.B. and Caskey,C.Thomas.
Oligonucleotide repeat arrays
Patent: US 5981185-A 56 09 NOV-1999;
Location/Qualifiers
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Similarity 89.5%;
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/mol_type="unassigned
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/mol_type="unassigned
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Pred. No. 97;
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AR242257
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Best Local S
Matches 17
                      Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 1.1%;
Best Local Similarity 89.5%;
Matches 17; Conservative
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Devices coated with substances that mediate
biological material
Patent: WO 2004055153-A 4 01-JUL-2004;
Eberhard-Karls-Universitaet Tuebingen (DE)
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Sequence 4 from Patent WO2004055153.
CQ830492
CQ830492.1 GI:50250832
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17; Conserv
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AR242257
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Ford, J. and Yeung, G.
Chemokine receptor obtained from a cDNA library of fetal
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llarity 89.5%;
Conservative
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                                     1.1%;
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                         Score 15.8; D
Pred. No. 97;
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AR242262
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Degussa Bioactives GmbH (DE)
Location/Qualifiers
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artificial sequences.
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Ford, J. and Yeung, G.
Chemokine receptor obtained from a cDNA library of fetal
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Location/Qualifiers
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/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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RESULT 53
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Sequence 7 from Patent WO03072818.
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synthetic construct synthetic construct
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/mol type="unassigned DNA"
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Location/Qualifiers
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Sequenz:Capture-Oligonukleotid"
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Sequenz:Capture-Oligonukleotid"
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|mol_type="unassigned DNA"
|db_xref="taxon:32630"
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/mod_base=OTHER
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                 note="LNA-T (Locked Nucleic Acid)"
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                                                              note="LNA-T (Locked Nucleic Acid)"
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Location/Qualifiers
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AX825112
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/mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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mod_base=OTHER
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Pred. No. 97;
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AX825115/c
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AX825113/c
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Sequence 11 from Patent WO03072818.
AX825113
AX825113.1 GI:39750842
                                                                    Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucle Patent: WO 03072818-A 13 04-SEP-2003; Degussa Bioactives GmbH (DE)
Location/Qualifiers
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synthetic construct
artificial sequences.
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Similarity 89.5%; Pred. No. 97;
17; Conservative 0; Mismatches
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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AX825116/c
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Sequence 14 from Patent WO03072818.
AX825116
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Sequenz:Capture-Oligonukleotid"
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/mb.tryce="bung der kuenstlichen
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AX825152/c
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AX825117/c
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synthetic construct synthetic construct artificial sequences.
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Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucle Patent: WO 03072818-A IS 04-SEP-2003; Degussa Bioactives GmbH (DE)
Location/Qualifiers
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Sequenz:Capture-Oligonukleotid"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/nod_base=OTHER
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89.5%; Pred. No. 97;
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AX825153
AX825153.1 GI:39750882
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Location/Qualifiers
                                                                                                                                                                                                                         Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids Patent: WO 03072818-A 51 04-SEP-2003; Degussa Bioactives GmbH (DE)
Location/Qualifiers
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/mol type="unassigned DNA"
/mol type="taxon:32630"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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                                                                           'note="LNA-T (Locked Nucleic Acid)"
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                               note="LNA-T (Locked Nucleic Acid)"
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AX825163/c
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Sequence 61 from Patent WO03072818.
AX825163
AX825163.1 GI:39750892
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|mol type="unassigned DNA"
|db xref="taxon:3530"
|noce="Beschreibung der kuenstlichen
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Sequence 19
AR164336
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I31828
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Torrence, P.F., Silverman, R.H., Maitra, R.K. and Lesiak, K. Chimeric molecules targeted to viral RNAs
Patent: US 6271369-A 19 07-AUG-2001;
Unclassified.
1 (bases 1 to 22)
Torrence, P., Silverman, R.,
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Methods for identification of cancer cell surface molecules cancer specific promoters, and therapeutic uses thereof Patent: WO 03000928-A 415 03-JAN-2003;
Odin Medical A/S (DK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (human)
Homo sapiens
Eukaryota, Metazoa; Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates; Catarrhini; Hominidae; Homo.
                                                Unknown.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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                                                                         GI:1822619
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S 5583032.
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    Maitra, R. and Lesiak, K.
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Sequence 19 from patent US 5677289.
169425
169425.1 GI:2831547
                                                                                                         PN JP 2002541795-A/2217

PD 10-DEC-2002

PD 11-APR-2000 JP 2000611654

PF 11-APR-2000 UP 2000611654

PR 12-APR-1999 US 60/129390

PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN PC C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC
                                                                                                                                                                                                                                              1 (bases 1 to 17)

Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.

Regulation of repressor genes using nucleic a

Patent: JP 2002541195-A 2217 10-DEC-2002;

RIBOZYME PHARMACEUTICALS INC
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1 (bases 1 to 22)

Torrence,P., Silverman,R., Maitra,R. and Lesiak,K.

Method of cleaving specific strands of RNA and medical treatments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Method of cleaving specific strands of Patent: US 5583032-A 19 10-DEC-1996; Location/Qualifiers
                                  C12P21/02, C12P21/02//A61K31/711, (C12N5/10, C12R1:91), (C12P21/02, PC C12R1:91),
                                                                                                                                                                                                                                                                                                                                                                     unidentified
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81.8%; Pred. No. 1e+02;
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Beigelman, L., Burgin, A., Beaudry, A., Karpeis
Matulic-Adamic, J., Sweedler, D. and Zinnen, S.
Synthetic ribonucleic acids with RNAse activ
Patent: US 6528640-A 559 04-MAR-2003;
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Key
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 2546 12-FEB-2002;
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559 from patent US 6528640.
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/mol_type="unassions
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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                                              /organism="unknown"
/mol_type="unassigned RNA"
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AR398177
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Sequence
AR323668
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Beigelman, L., Burgin, A.B., Beaudry, A., Karpeisky, A., Matulic-Adamic, J., Sweedler, D. and Zinnen, S. Oligoribonucleotides with enzymatic activity Patent: US 6617438-A 558 09-SEP-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 17)
Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6566127-A 1070 20-MAY-2003;
Location/Qualifiers
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Oreochromis niloticus (Nile tilapia)
Oreochromis niloticus
Eukaryota; Metazoa; Chordata; Crania
                                                                                                                                                                                                   16;
                                                                                                                                                                                                                                                                                                                                                                         Unknown
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Similarity 94.1%; Pred. No. 1.5e+02;
16; Conmervative 0; Mismatches 1
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/mol_type="unassigned RNA"
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/mol_type="unassign
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94.1%; Pred. No. 1.5e+02;
tive 0; Mismatches 1
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  Chordata; Craniata; Vertebrata;
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Sequence
A90272
                                                    l (bases 1 to 20)
Brysch, W.D. and Schlingensiepen, K.D.
An antisense oligonucleotide preparation
Patent: EP 0856579-A 453 05-AUG-1998;
BIOGNOSTIK GES (DE)
                                                                                                                     unidentified unclassified.
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Brysch, W. and Schlingensiepen, K.

Brysch, W. and Schlingensiepen, K.

ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD Patent: WO 9833904-A 453 06-AUG-1998;

BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           unidentified unidentified
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Sequence 453 from Patent
A88305
A88305.1 GI:6736875
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Labroidei; Cichlidae; Oreochromis.
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
/db_xref="taxon:8128"
                                         Location/Qualifiers
/organism="unidentified"
/mol_type="unassigned DNA"
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94.1%; Pred. No. 1.3e+02;
tive 0; Mismatches 1;
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AUTHORS
AN antisense oligonucleotide preparation method
FITLE
FACENCY A 43 07-AUG-2001;
BIGGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
OS Unknown
OS Unknown
OS Unknown
OF AUG-2001
PD 07-AUG-2001
PF 30-JAN-1998 JP 1998532533
PD 07-AUG-2001
PF 30-JAN-1998 JP 1998532533
PI KARL HERMANN SCTT
PC C12N15/**
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AX356277
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An antisense oligonucleotide
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Plant regulatory sequences for selective control of gene expression
Patent: WO 0200905-A 71 03-JAN-2002;
Monsanto Technology LLC (US)
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Similarity 94.1%;
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                                                     /organism='Unknown'.
/organism="unidentified"
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Pred. No. 1.3e+02;
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Pred. No. 1.3e+02;
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AR112167/c
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Best Local Similarity
Matches 16; Consert
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AR112167 20 bp
Sequence 56 from patent US 6130041.
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Acton, S. Laurene.
Intronic and polymorphic SR-BI nucleic acids and Patent: US 5998141-A 56 07-DEC-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                 Sequence 56 from patent US AR092032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genotyping reagents, kits and methods of use thereof Patent: WO 0129262-A 3390 26-APR-2001; Orchid BioSciences, Inc. (US)
                                                                                                                                                                                                                                                                                                Unknown.
Unclassified.
                                                                                                                                                                                                                                                                                                                           Unknown.
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Similarity 94.1%;
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ilarity 94.1%;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Primer"
                                                                                                                                                                                    /mol_type="unassigned DNA"
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Pred. No. 1.2e+02;
0; Mismatches 1;
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Pred. No. 1.
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BD229218
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AR112167.1
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (base 1 to 20)

Galvin,M., Miller,A., Penny,L. and Riedy,M.
Genotype determination of human UDP-glucuronosyl transferase 2B4 (UGT2B4), 2B7 (UGT2B7) and 2B15 (UGT2B15) genes
Patent: JP 2002521067-A 90 16-JUL-2002;

AXYS PHARMACEUTICALS INC
OS Homo sapiens (human)
                                                                                                                                                              Genotype determination (UGT2B4), 2B7 (UGT2B7) & BD229218 BD229218.1 GI:33038988
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Acton, S.L. and Ordo
Human intronic and
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Sequence 56 from patent US
AR149209
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Acton, S. Laurene.
Human intronic and polymorphic SR-BI nucleic acids and uses
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Location/Qualifiers
                                                                                                                          Homo sapiens
                                                                                                                                     Homo sapiens (human)
                                                                                                                                                    JP 2002521067-A/90.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polymorphic SR-BI nucleic
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; Pred. No. 1.5e
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UDP-glucuronosyl transferase
(UGT2B15) genes.
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AR349470
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Best Local Similarity
Matches 17; Conserv
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Best Local Similarity 85.0%;
Matches 17; Conservative
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PD 16-JUL-1902

PF 22-JUL-1999 JP 2000562558

PR 28-JUL-1999 US 60/094391

PI MARGARET GALVIN, ANDREW MILLER, LAURA PENNY, MICHAEL RIEDY F

C12N15/09, C12N15/09, C12M1/00, C12Q1/68, C12N15/00, C12N15/00 CC

Genotype determination of human UDP-glucuronosyl transferase

284 (UGT2B4),

294 (UGT2B4) (UGT2B15) genes
                                   1 (bases 1 to 20)
Galvin, M., Miller, A., Penny, L. and Riedy, M.
Genotyping the human UDP-glucuronosyltransferase
Patent: US 6586175-A 92 01-JUL-2003;
Location/Qualifiers
                                                                                                                    Unknown
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Sequence 92 from patent US
AR349470
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Aizawa,A., Kawakami,A. and Kondo,T.
Testis-specific gene
Patent: US 6555670-A 4 29-APR-2003;
Location/Qualifiers
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Sequence 4 from patent US
AR309844
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llarity 85.0%;
Conservative
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic"
             organism="unknown"
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_type="genomic DNA"
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Pred. No. 1.5e+02;
0; Mismatches 3
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Mus sp.
Eukaryota; !
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity 85.0
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17; Conserv
                                                                                                                                                                    Livestock Improvement Association of Gunma University (JP)
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                                                                                                                                                                                             Testis-specific gene Patent: EP 1195382-A 4 10-APR-2002;
                                                                                                                                                                                                                                                           synthetic construct synthetic construct
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AX298541
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                   AATCTATTTTTTTATTTTC 1319
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AAGCTTTTTTTTTTTTTTTC
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                                                  Conservative
                                                                                               /organism="synthetic construct"
/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus sp."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                     Location/Qualifiers
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                                                           1.1%;
85.0%;
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Pred. No. 1.5e+02
0; Mismatches
                                               Score 15.2; DB 1
Pred. No. 1.5e+02
0; Mismatches
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Pred. No. 1.5e+02
0; Mismatches
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                                                                                                                                                                                  (JP) ; President
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SOURCE
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                                                                                    RESULT 88
BD143136
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AUTHORS
TITLE
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AX770111
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Matches 17
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Best Local Similarity
Matches 17; Conserv
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AX488408
             28D143136
Novel testis-specific gene.
BD143136
BD143136.1 GI:27848894
JP 2002112777-A/3.
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                                                                                                                                                                                                      l Similarity
17; Conserv
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Sequence 9 from Patent WO03016562.
AX770111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       synthetic construct synthetic construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AX770111.1 GI:32437689
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ilarity 85.0%;
Conservative
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llarity 85.0%;
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/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                /mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Primer"
                                                                                                                                                                                                                                                                                                        organism="synthetic construct"
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Pred. No. 1.5e+02;
0; Mismatches 3
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Pred. No. 1.5e+02;
0; Mismatches 3
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AX825106
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       synthetic construct synthetic construct artificial sequences.
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Sequence 4 from Patent WO03072818.
AX825106
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1 (bases 1 to 20)

Aizawa,A., Kawakami,A. and Kondo,T.

Novel testis-specific gene
Patent: JP 2002112777-A 3 16-APR-2002;

KACHIKU KAIRYO JIGYODAN, PRESIDENT OF GUNMA UNIVERSITY
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Similarity 85.0%; Pred. No. 1.5e+02
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AKIRA AIZAWA, AKIKO KAWAKAMI, TOSHIHIKO
C12N15/09.C07K14/47,C12N15/00
Novel testis-specific gene
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JP 2002112777-A/3
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                                                                                                                                                                                                                                                                          /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="texon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism='Artificial
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                                                      mod_base=OTHER
                                                                                        note="LNA-T (Locked Nucleic Acid)"
                                                                                                                note="LNA-T (Locked
mod_base=OTHER
                                                                                                                                                           /note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
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Matches 17; Conserv
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Local Similarity 85.0%;
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A20531 21 bp D coligonucleotide for the mutagenesis
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17; Conserv
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synthetic construct
artificial sequences.
1 (bases 1 to 21)
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synthetic construct
artificial sequences.
1 (bases 1 to 21)
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Location/Qualifiers
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milarity 85.0%;
Conservative 0
                                                                                                                           1.1%;
larity 85.0%;
Conservative
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/db_xref="taxon:32630"
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Pred. No. 1.4e+02;
0; Mismatches 3
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Pred. No. 1.4e+02;
0; Mismatches 3;
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AR143626
AR143626.1 GI:15104912
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A20531.1 GI:583363
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Location/Qualifiers
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             Patent: US 6204435-A 11 20-MAR-2001;
Location/Qualifiers
                                                  Stamp, L.M.
Pesticidal toxins and nucleotide sequences which encode these
                                                                    1 (bases 1 to 21)
Feitelson,J.S., Schnepf,H.Ernest., Narva,K.E., Stockhoff,B.A., Schmeits,J., Loewer,D., Dullum,C.Joseph., Muller-Cohn,J. and Stamp 1. M
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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I (bases) to 21)

Babba, T. and Shimada, H.

NEW RICE PLANT STARCH-BRANCHED ENZYMIC GENE
PATENT: JP 1994261767-A $ 20-SEP-1994;

MITSUI GIYOUSAI SHOKUBUTSU BIO KENKYUSHO:KK
OS None
OC Attificial sequences.
PN JP 1994261767-A/5
PD 20-SEP-1994
PF 22-OCT-1993 JP 1993265171
PR 29-OCT-1993 JP 1993265171
PR 20-OCT-1993 JP 1993265171
PR 20-
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Feitelson, J.S., Schnepf, H. Ernest., Narva, K.E., Stockhoff, B.A., Schmeits, J., Loewer, D., Dullum, C. Joseph., Muller-Cohn, J., Stam Morrill, G. and Finstad-Lee, S.
Pesticidal toxins and nucleotide sequences which encode these
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JP 1994261767-A/5.
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Location/Qualifiers
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Artificial sequences.

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20-SEP-1994
22-OCT-1993 JP 1993265171
29-OCT-1993 JP 1993265171
29-OCT-1992 JP 92P 291719
EABA TADASHI, SHIWADA HIROAKI
C12N15/54,AO1H5/00,C12N5/10,C12P19/16//A23L1/10,C12N9/10;
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/mol_type="unassigned DNA"
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    /organism='Artificial sequences'.
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Pred. No. 1.4e+02;
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AR437180
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Location/Qualifiers
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Feitelson, J.S., Schnepf, H.E., Narva, K.E., Stockhoff, B.A., Schmeits, J., Loewer, D., Dullum, C.J., Muller-Cohn, J., Stamp, L., Morrill, G. and Finstad-Lee, S.
Pesticidal toxins and nucleotide sequences which encode these
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Location/Qualifiers
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/db xref="taxxon:32630"
/noTe="Beschreibung der kuenstlichen
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/note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
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from patent US 6656908.
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85.0%; Pred. No. 1.4e+02;
tive 0; Mismatches 3;
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/mod_base=OTHER
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/mod_base=OTHER
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/mod_base=OTHER
                                                                                                                                                                                                                                                                                                                                                                                                organism="synthetic construct"
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/db_xref="taxon;32630"
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/mod_base=OTHER
                                                                              'note="LNA-T (Locked Nucleic Acid)"
'mod_base=OTHER
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Pred. No. 1.4e+02;
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Sequence 33
AX825135
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AX825121
          Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucle Patent: WO 03072818-A 33 04-SEP-2003; Degussa Bioactives GmbH (DE)

Location/Qualifiers
                                                                                                            synthetic construct synthetic construct
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/mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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Pred. No. 1.4e+02;
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Sequence 34 from Patent WO03072818.
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17; Conserv
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sequenz:Capture-Oligonukleotid"
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TITLE
                                                      RESULT 104
AX825137/c
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KEYWORDS
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/mod_base=OTHER
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FEATURES

SOURCE

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21 bp
Sequence 34 from Patent WO03072818.
AX825136
AX825136.1 GI:39750865
AX825137 21 bp
Sequence 35 from Patent W003072818.
AX825137 GI:39750866
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Location/Qualifiers
                                                                                                                                                                                                  1.1%; Score 15.2; DB 1; larity 85.0%; Pred. No. 1.4e+02; Conservative 0; Mismatches 3;
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Sequenz:Capture-Oligonukleotid"
                                                                                                                                                                                                                                                                 /note="LNA-T (Locked Nucleic
/mod_base=OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="synthetic construct"
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/db_xref="taxon:32630"
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mod_base=OTHER
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Pred. No. 1.4e+02;
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Sequence 53 from Patent WO03072818.
AX825155
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Sequenz:Capture-Oligonukleotid"
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/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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                              note="LNA-T (Locked Nucleic Acid)"
                                                         bound_moiety="Biotin"
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Pred. No. 1.4e+02;
0; Mismatches 3;
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TTAAAAAAAAAAAGTAAAA 1537
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21 bp
Sequence 57 from Patent WO03072818.
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Location/Qualifiers
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 Conservative
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/mod_base=OTHER
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/mod_base=OTHER
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/mol_type="unassigned DNA"
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/noTe="Beschreibung der kuenstlichen
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/mod_base=OTHER
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mod_base=OTHER
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/mod_base=OTHER
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               1.1%; Score 15.2; DB 1;
85.0%; Pred. No. 1.4e+02;
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85.0%; Pred. No. 1.4e+02;
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hes 17; Conserv
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                                                                                                                  BD023735

Beta-galactosidase having reversibly inactive lactase activity.
BD023735

BD023735.1 GI:22564958

JP 2001506136-A/1.
Exemothecium gossypii (Ashbya gossypii)
Eremothecium gossypii Bremothecium gossypii Bremothecium gossypii Bremothecium gossypii Saccharomycetae; Saccharomycetaes; Saccharomycetaes; Saccharomycetaes;
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PI JAMES L SCHMEITS, DAVID LOEWER, GEORGE SCHWAB,

PI CHARLES JOSEPH DULLUM,

PI JUDY MULLER COHN, LISA STAMP

PC C12N15/32,C07K14/325,C12Q1/68,A01N63/00,C12N15/82

Strandedness: Single;

CC Topology: Linear;

FH Key

Location/Qualifiers

FH Key

1. 21

FT source

/organism='Unidentified'.
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Novel pesticidal toxins and nucleotide sequences which encode these
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Peitelson,J.S., Schmepf,E.H., Narva,K.E., Stockhoff,B.A.,
Schmeits,J.L., Loewer,D., Schwab,G., Dullum,C.J., Cohn,J.
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JP 2001502919-A/8.
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JP 2001502919-A/8
06-MAR-2001
30-OCT-1997 JP 1998520788
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pesticidal toxins and nucleotide sequences which encode these
                                                                                                                                                                                                                                                                                                                                                                                        /organism="unidentified"
/mol_type="genomic DNA"
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Pred. No. 1.4e+02;
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AX117632
LOCUS
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SOURCE
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AR187059/c
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Sequence 2547 f
AR187059
AR187059.1 GI:
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1511 CTGTTAATTAAAAAAAAAA 1530
                                              1 (bases 1 to 17)
Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 2547 12-FEB-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Picoult-Newburg,L. and Pohl,M. Genotyping reagents, kits and methods . Patent: WO 0129262-A 2755 26-APR-2001; Orchid BioSciences, Inc. (US)
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Strandedness: Single;
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/mol_type="genomic DNA"
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/mol_type="unassigned
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REFERENCE
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TITLE
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AR323669/c
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AR323670/c
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AR187060/c
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Best Local Similarity 100.0%; Pred. No. 1.9e+02;
Matches 15; Conservative 0: Mismatches 0.
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Sequence 2548 from patent
AR187060
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Sequence 1071 from patent
AR323669
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1 (bases 1 to 17)

Pavers, McSwiggen, J., Stinchcomb, D. and Escobedo, J.

Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398 A 2548 12-FEB-2002;
         AR323670
Sequence 1072 from patent
AR323670
                                                                                                                                                                                                                                    1 (bases 1 to 17)
Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6566127-A 1071 20-MAY-2003;
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 AR323670.1
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ilarity 100.0%;
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 GI:33709478
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100.0%; Pred. No.
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                                     PAT 17-AUG-2003
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AX753821/c
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FEATURES
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KEYWORDS
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                                                                                                                                                  Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primatos;
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Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6566127-A 1072 20-MAY-2003;
Location/Qualifiers
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                    Sequence 169 from Patent AX753822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shannon,M. and Phan,T.
Human angiomotin-like protein 1
Patent: WO 03037931-A 168 08-MAY-2003;
Amersham Biosciences SV Corp. (US)
Location/Qualifiers
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                                  Human angiomotin-like protein 1
Patent: WO 03037931-A 169 08-MAY-2003;
Amersham Biosciences SV Corp. (US)
Location/Qualifiers
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                                                                                                                  Shannon, M. and Phan, T.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned RNA"
/organism="Homo sapiens"
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tive 0; Mismatches 0
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Sequence 170 from Patent WO03037931.
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Sequence 18 from patent US 6083731.
AR102020
AR102020.1 GI:12812818
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                                                                                                                                                                                                  Unknown.
Unclassified.
1 (bases 1 to 19)
1 (bases 1 to 19)
Croteau, R. Bruce., Lupien, S. Lee. and Karp, F.
Recombinant materials and methods for the p
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae, Homo.
                                                                                                                                                                  Patent: US 6083731-A 18 04-JUL-2000; Location/Qualifiers
                                                                                                                                                                                                                                                                 Unknown.
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Patent: WO 03037931-A 170 08-MAY-2003;
Amersham Biosciences SV Corp. (US)
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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tive 0; Mismatches 0;
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AR352460/c
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JP 1999196874-A/9
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1 (bases 1 to 20)
Hideki,K. and Senshu,U.
Method for analyzing DNA fragment
Patent: JP 1999196874-A 9 27-JUL-1999;
                                                                                                                  E28098
E28098.1 GI:13018323
JP 1999196874-A/9.
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Method for analyzing DNA fragment.
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Kong,Y., Chung,J.-Y., Bahk,Y.Y., Kang,S.-Y. and Cho,
Recombinant antigen of Taenia solium metacestodes
Patent: US 6589752-A 6 08-JUL-2003;
Location/Qualifiers
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Croteau, R. Bruce., Lupien, S. Lee, and Karp, F.
Recombinant materials and methods for produced
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Similarity 84.2%; Pred. No. 1.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                    Unclassified.
1 (bases 1 to 20)
1 (bases 1 to 20)
Cohen,D., Chumakov,I. and Blumenfeld,M. Biallelic markers for use in constructing disequilibrium map of the human genome patent: US 6537751-A 6435 25-MAR-2003;
Location/Qualifiers
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PP 14-JAN-1998 JP 1998005399
PR HIDEKI KAMIBARA, SENSHU UEMATSU
PI C12N15/09, C12Q1/68, G01N27/447, C12N15/00, G01N27/26
Strandedness: Single;
CC Topology: Linear;
FH Key Location/Qualifiers
FH Key 1...20
FT Bource 1...20
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                                                                                                                                                             AR307942
Sequence 153 from patent
AR307942
AR307942.1 GI:31698698
                                      Antisense modulation of raidd expression Patent: US 6551826-A 153 22-APR-2003; Location/Qualifiers
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Sequence 6435 from patent
AR294700
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/db_xref="taxon:32644"
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Location/Qualifiers
              organism="unknown"
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mol_type="genomic DNA"
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84.2%; Pred. No. 1.6e+02;
Live 1; Mismatches 2;
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100.0%; Pred. No. 1.6e+02;
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AX048446/c
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VERSION
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AX048446
AX048446.1
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Detection system for separating constituents
production and use of the same
Patent: WO 0071747-A 45 30-NOV-2000;
Aventis Research & Technologies GmbH & Co. KC
Location/Qualifiers
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                                                                                                                                                                                                                                                                                              Oreochromis niloticus (Nile tilapia) Oreochromis niloticus
                                                                                                                                                                                                                                                                                                                                                 AX804844
Sequence 1012 from Patent
AX804844
                                                                                                                                                              Genomar ASA (NO)
                                                                                                                                                                                                  Lie,Y., Slettan,A., Hoeyum,M. and Lingaas,F. Verification of food origin based on nucleic
                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygi; Neopterygi; Teleostei; Euteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Labroidei; Cichlidae; Oreochromis.
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                                                                                      /organism="Oreochromis niloticus"
/mol_type="unassigned DNA"
/db_xref="taxon:8128"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequenz:Erkennungssystem"
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/db_xref="taxon:32630"
/note="Beschreibung der kunstlichen
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AR058305
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Best Local Similarity 88.9%;
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Sequence 3 :
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Chatelain, F. and Kumarev, V.

Process for preparing polynucleotides on tightly packed bed 
Patent: US 5869643-A 18 09-FEB-1999;

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l (bases 1 to 18)
Chatelain, F. and Kumarev, V.
Process for preparing polynucleotides on tightly packed bed
Patent: US 5869643, A 12 09-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
AR034896
Unclassified.
1 (bases 1 to 18)
De Rose,R., Douce,R.,
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Sequence 18 from patent US
AR034899
AR034899.1 GI:5950504
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                                                                                 from patent US
                                                               GI:5983882
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Pred. No. 2e+02;
  Duval,M.,
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Unknown.
                                                          16;
                                                                                                                                                               Unclassified.

1 (bases 1 to 18)

Manly,S.P., Kozlowski,M.R. and Neve,R.L.

Cloning and expression of .beta.APP-C100

Patent: US 6083713-A 25 04-JUL-2000;

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                      AR101834
Sequence 25 from patent US
AR101834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            biology experiments
biology experiments
Patent: US 6071745-A 9 06-JUN-2000;
Location/Qualifiers
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Lin,C.-I.Patsy., Wallace,R.Bruce, Cossman,J. and French,C.
Method and formulation for lyophilizing cultured human cells to
preserve RNA and DNA contained in cells for use in molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unknown
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AR097579
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Similarity 88.9%; Pred. No. 2e+02;
16; Conservative 0; Mismatches 2;
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/mol_type="unassigne
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/mol_type="unassigned DNA"
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/mol_type="unassigned
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88.9%; Pred. No. 2e+1
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KEYWORDS
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BD222596/c
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AUTHORS
TITLE
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FEATURES
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Best Local Similarity 88.9%;
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Cowsert,L.M. Bennett,C.Frank. and O'Malley,B.W.
Antisense modulation of SRA expression
Patent: US 6107092-A 70 22-AUG-2000;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AR106909
Sequence 70 from patent US
AR106909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AR106506
Sequence 30 from patent US
AR106506
produced therefrom
Patent: JP 2002522447-A 14 23-JUL-2002;
                        synthetic construct synthetic construct artificial sequences. artificial sequences. 1 (bases 1 to 18) 1 (bases 1 to 08) Manoharan, M., Cook, P.D., Prakash, T.P. and Kawasaki, A.M. Aminooxy-modified nucleoside compound and oligomer compound
                                                                                                        produced therefrom.
BD222596
BD222596.1 GI:33032366
JP 2002522447-A/14.
                                                                                                                                                              BD222596 18
Aminooxy-modified nucleoside
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unknown.
Unclassified.
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Patent: US 6107060-A 30 22-AUG-2000;
Location/Qualifiers
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Keeling, P. and Guan, H.
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                                                                                                                                                                                                                                                                                                                                                         /mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                                                         organism="unknown"
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Pred. No. 2e+02;
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Pred. No. 2e+02;
0; Mismatches 2;
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S 6107092.
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Best Local Similarity 88.9%;
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                          1520 AAAAAAAAAAAGTAAAA 1537
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Kenichi, H., Hiroshi, Y. and Masahide, N.
Method for labeling oligonucleotide and utilization thereof
Patent: JP 199075880-A 2 23-MAR-1999;
CHEMO SERO THERAPEUT RES INST
OS Unidentified
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                                                                                                                                                                                                                         PI KENICHI HANAKI, HIROSHI YOSHIKURA, MASAHIDE NOZAKI
C12N15/09, C12Q1/68, G01N33/58, C12N15/00
CC Strandedness: Single;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E28535
Method for labeling
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                                                                n 1.1%;
Similarity 88.9%;
 AAAAAAAAAAAAAAAAA 18
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JP 199075880-A/2
23-MAR-1999
10-JUL-1998 JP 1998195719
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07-AUG-1998 US 09/130973
MUTHIAH MANOHARAN, PHILIP DAN COOK, THAZHA P PRAKASH, ANDREW M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Artificial Sequence JP 2002522447-A/14 23-JUL-2002
                                                    Conservative
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C12N15/00
                                                                                                                                                                                                             Topology: Linear;
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                                                                                                    /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                                      /organism='Unidentified'.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism='Artificial Sequence'
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oligonucleotide
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                                                Score 14.8; DB 1;
Pred. No. 2e+02;
0; Mismatches 2;
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Pred. No. 2e+02;
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RESULT 134 E28536/c

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AUTHORS
TITLE
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AR196702/c
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179509/c
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PN JP 199075880-A/3
PD 23-MAR-1999
PF 10-JUL-1998 JP 1998195719
PR KENICHI HANAKI, HIROSHI YOSHIKURA, MASAHIDE NOZ
C12N15/09, C12Q1/68, GQ1N33/58, C12N15/00
CC Strandedness: Single;
CC Topology: Linear;
FH Key Location/Qualifiers
FT source 1.18
FT source /organism='Unidentified'.
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Kenichi,H., Hiroshi,Y. and Masahide,N.

Method for labeling oligonucleotide and utilization thereof

Patent: JP 1999075880-A 3 23-MAR-1999;

CHEMO SERO THERAPEUT RES INST
                                                                                                                                                                                                                                                                                                                                                                        Sequence 16
179509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism='Unidentified'
                                                                                                                                                                                                 /mol_type="unassigned DNA"
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Pred. No. 2e+02;
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5 5707807.
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2e+02;
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AR196704/c
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AR215435/c
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Matches 16
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                                                                                                                                                                                                 Sequence 9 AR215435
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                                       i (bases 1 to 18)
Lin,C.-I.P., Wallace,R.B., Cossman,J. and French,C.
Method and formulation for lyophilizing cultured human cells to
preserve RNA and DNA contained in cells for use in molecular
biology experiments
biology experiments
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 18)
Zwick, M.G., Edington, B.E., McSwiggen, J.A., Mc
Guo, L., Skokut, T.A., Young, S.A., Folkerts, O.
Nucleic acid encoding delta-9 desaturase
Patent: US 6350934-A 1169 26-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1169 from patent AR196704 AR196704.1 GI:20246141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Zwick, M.G., Edington, B.E., McSwiggen, J.A., Merlo, P.Ann. Owens., Guo, L., Skokut, T.A., Young, S.A., Folkerts, O. and Merlo, D.J. Nucleic acid encoding delta-9 desaturase Patent: US 6350934-A 1167 26-FEB-2002;
                                                                                                                                                                                                                             AR215435
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ilarity 88.9%;
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/mol_type="unassigned DNA"
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/organism="unknown"
/mol_type="genomic |
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D. and Merlo, D.J.
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PAT 20-APR-2002

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25-SEP-2002

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AR473365/c
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AR412363/c
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AR473365
Sequence 9 :
AR473365
AR473365.1
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Sequence 24 from patent US
AR222464 AR222464.1 GI:23329995
                                                                                                                                                                                                                       1 (bases 1 to 18)
Manoharan, M., Cook, P.D., Prakash, T.P. and Kawasaki, A.M.
Aminooxy-modified nucleosidic compounds and oligomeric
prepared therefrom
Patent: US 6639062-A 14 28-OCT-2003;
                                                                                                                                                                                                                                                                                                                                   AR412363
Sequence 14 from patent US
AR412363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unclassified.
1 (bases 1 to 18)
Kurz,M., Lohse,P. and Wagner,R.
Kurz,M., Lohse,P. and Wagner,R.
Peptide acceptor ligation methods
Patent: US 6429300-A 24 06-AUG-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                        Unknown.
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                                                                                                                                                                                                                                                                               Unclassified.
                                                                                                                                                                                                                                                                                                   Unknown.
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/mol_type="genomic DNA"
                                                                                                                                                                                                              Location/Qualifiers
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/mol_type="genomic DNA"
                      from patent US
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  GI:42708816
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Pred. No. 2e+0
0; Mismatches
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Pred. No. 2e+02;
D; Mismatches
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AR487020/c
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AR487019
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Sequence 7
AR487020
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Sequence 6
AR487019
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Lin,C.-I.P., Wallace,R.B., Cossman,J. and French,C.
Method and formulation for lyophilizing cultured hu
preserve RNA and DNA contained in cells for use in
biology experiments
Patent: US 6686460-A 9 03-FEB-2004;
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                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 18)
Thirstrup,K., Warthoe,P. and Pettersson,N.B.
Process for amplifying and labeling single s
ligated adaptor mediated amplification
Patent: US 6706476-A 6 16-MAR-2004;
Location/Qualifiers
                                    1 (bases 1 to 18)
Thirstrup, K., Warthoe, P. and Pettersson, N.B
Process for amplifying and labeling single
ligated adaptor mediated amplification
Patent: US 6706476-A 7 16-MAR-2004;
Location/Qualifiers
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Unclassified
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/organism="unknown"
/mol_type="genomic |
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; Pred. No. 2e+0
0; Mismatches
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Pred. No. 2e+02;
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Best Local Similarity 88.9%;
Matches 16; Conservative
                                                      Matches
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Method for isolating anionic organic substances from
systems using cationic polymer nanoparticles
Patent: WO 9910527-A 4 04-MAR-1999;
SUEDDEUTSCHE KALKSTICKSTOFF (DE); BAYER ERNST (DE)
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Sequence 4 from Patent WO9910527.
AXOO4875
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Method for isolating anionic organic substances fr
systems using cationic polymer nanoparticles
patent: WO 9910527-A 8 04-MAR-1999;
SUEDDEUTSCHE KALKSTICKSTOFF (DE); BAYER ERNST (DE)
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Sequence 8 from Patent WO9910527.
AX004879
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16; Conserv
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/mod_base=um
                                                                                                                                     /organism="synthetic construct"
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/note="2' methyl-modified oligonucleotide"
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                                                                   Score 14.8; DB 
Pred. No. 2e+02;
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                        1520 AAAAAAAAAAAGTAAAA 1537
18
                                                                                                                                                                                       and its analogues
Patent: WO 9967378-A 3 29-DEC-1999;
DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL A (CA); WILDS CHRISTOPHER
DAMHA MASSAD JOSE (CA); PARNION DOMINIQUE (CA); NORONHA ANNE M (CA)
(CA); UNIV MCGILL (CA); ARION DOMINIQUE (CA);
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Patent: WO 9967378-A 2 29-DEC-1999;
DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL
(CA); UNIV_YONE (CA); ARION DOMINIQUE
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AX008117
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llarity 88.9%;
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88.9%; Pred. No. 2e+02;
tive 0; Mismatches 2;
                                                    Score 14.8; D
Pred. No. 2e+0
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RESULT 148

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AX028843
Sequence 27 from Patent WO9732023.
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Patent: WO 9967378-A 8 29-DEC-1999;
DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL
(CA); UNIV_MCGILL (CA); ARION DOMINIQUE
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Sequence 8 from Patent WO9967378.
AX008123
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Sequence 7 from Patent WO9967378.
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Patent: WO 9967378-A 7 29-DEC-1999;

DAWHA MASSAD JOSE (CA); PARNIAK MICHAEL A (CA); WILDS CHRISTOPHER

(CA); UNIV MCGILL (CA); ARION DOMINIQUE (CA); NORONHA ANNE M (CA);
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88.9%; Pred. No. 2e+02;
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Gumbiowski, K. and Zulauf, M.
High density labeling of dna with modified or
nucleotides and dna polymerases used
Patent: WO 0068422-A 21 16-NOV-2000;
Roche Diagnostics GmbH (DE)
Location/Qualifiers
                                                                                          Sequence 23 from Patent WO0068422. AX047273
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Sequence 21 from Patent WO0068422.
AX047271
synthetic construct artificial sequences.
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FLORIGENE LIMITED (AU); BRUGLIERA FILIPPA
ALBERT (AU); MICHAEL MICHAEL ZENON (AU)
Location/Qualifiers
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Genetic sequences encoding flavonoid pathway enzymes and uses
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Similarity 88.9%; Pred. No. 2e+02;
16; Conservative 0; Mismatches 2.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db xrefs"ttaxon:32630"
/noTe="Oligonuclectide"
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Pred. No. 2e+02;
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AX104747/c
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High density labeling of dna with modified or chromophore carrying nucleotides and dna polymerases used patent: WO 0068422-A 23 16-NOV-2000;
Roche Diagnostics GmbH (DE)
                                                                                                                                                                                                             AX104747
Sequence 939 from Patent
AX104747
AX104747.1 GI:13920944
                                                                            Krieg, A.M., Schetter, C. and Vollmer, J.C. Immunostimulatory nucleic acids Patent: WO 0122972-A 939 05-APR-2001; UNIVERSITY OF IOWA RESEARCH FOUNDATION (
                                                                                                                                                   synthetic construct synthetic construct artificial sequences.
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Immunostimulatory nucleic acids
Patent: WO 0122972-A 913 05-APR-2001;
UNIVERSITY OF IOWA RESEARCH FOUNDATION (US) ; Coley Pharmaceutical
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Similarity 88.9%;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/wol_type="unassigned DNA"
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AX108642/c
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AX105651/c
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Sequence 10 from Patent WO0123564.
AXIO5651
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Differentially expressed genes
Patent: WO 0123419-A 10 05-APR-2001;
SCIOS INC. (US)
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Similarity 88.9%;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                                                                              /note="synthetic"
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Pred. No. 2e+02;
0; Mismatches
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Sequence 913 from Patent
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AX547774.1 GI:25812918
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Sequence 837 from Patent
AX355809
AX355809.1 GI:18620477
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Sequence 84 from Patent WO0174901.
AX268883
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UNIVERSITY OF IOWA RESEARCH FOUNDATION
Location/Qualifiers
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Similarity 88.9%;
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/mol_type="unassigned DNA"
/db_xref="taxon:132630"
/noTe="Oligos corresponding to polylinker sequence.
                                                                                                                                                                                                                     backbone"
                                                                                                                                                                                                                              /mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Synthetic oligonucleotide~phosphorothioate
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KEYWORDS
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AX547800/c
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Inhibition of angiogenesis by nucleic acids
Inhibition of angiogenesis by nucleic acids
Patent: WO 02053141-A 939 11-JUL-2002;
Coley Pharmaceutical Group, Inc. (US)
Location/Qualifiers
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Sequence 1 from Patent WO03064441.
AX814716
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Sequence 939 from Patent
AX547800
                                                                                              Damha, M.J. and Parniak, M.A. Oligonucleotides comprising alternating segments and uses thereof Patent: WO 03064441-A 1 07-AUG-2003; MCGILL UNIVERSITY (CA)
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/organism="synthetic construct"
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/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol type="unassigned DNA"
/pe="unassigned DNA"
/db xrefs"taxon:32630"
/noTe="Synthetic Sequence"
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16; Conserv
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Sequence 8 from Patent WO03064441.
AX814723
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Similarity 88.9%;
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                                                /note="Residues 1-3, 7-9,
2'-O-methyl-D-uridine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="synthetic construct"
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/db_xref="taxon:32630"
                                                                                               organism="synthetic construct"
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/db_xref="taxon:32630"
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           8; DB 1;
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Oligonucleotides comprising alternating Patent: WO 03064441-A 21 07-AUG-2003; MCGILL UNIVERSITY (CA)
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Sequence 21 from Patent WO03064441.
AX814736
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Similarity 88.9%; Pred. No. 2e
                                                                                                                                                                                                                /organism="synthetic construct"
/mol_type="unassigned RNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
                                                                                                                                                                                                     /note="Target RNA oligonucleotide"
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                                                                                                                                      Score 14.8; DB 1; Length 18; Pred. No. 2e+02; O; Mismatches 2; Indels
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UNITECH CO LTD
OS Homo sapiens (human)
PN JP 2001333800-A/2
PD 04-DEC-2001
PF 30-MAY-2000 JP 2000160324
PI KAORI SHIVADA
PC C1201/68,C12N15/09,G01N33/50,C12N15/00
CC Method of comparison and detection of RNA amoun
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A68209
AR048767
Sequence 1 from
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COllingwood, S.P., Moser, H.E., Altmann, K. and Douglas, INTERNEDIATES FOR OLIGONUCLECTIDE SYNTHESIS
Patent: WO 9747636-A 4 18-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                           unidentified unidentified
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16; Conserv
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JP 2001333800-A/2.
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                                                                                                                                                                                                                                                                                                                                              unclassified.
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ilarity 88.9%;
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4 from Patent WO9747636.
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/mol_type="genomic RNA"
/db_xref="taxon:9606"
                                                                                                                                                                                              /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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Pred. No. 1.9e+02;
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AR111371/c
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Sequence 1 :
AR111371
AR111371.1
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Leclerc,G. and Martel,R.
Radiolabeled DNA oligonucleotide and method
Patent: US 5821354-A 1 13-OCT-1998;
Location/Qualifiers
                                                                    1 (bases 1 to 19)
Cook, P.Dan., Mancharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 20 03-CCT-2000;
Location/Qualifiers
                                                                                                                                            Unknown.
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Sequence 20 from patent US
AR111946
AR111946.1 GI:12828794
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Leeds,J.M. and Cummins,L.L.
Fluorescence based nuclease assay
Patent: US 6127124-A 1 03-OCT-2000;
Location/Qualifiers
                                                                                                                                                        Unknown.
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                                 /organism="unknown"
/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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AR111949/c
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AR111948/c
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Sequence 21 from patent US 6127533.
AR111947.1 GI:12828795
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           Unknown
                       Unknown.
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 22 03-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                Unknown
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 21 03-OCT-2000;
Location/Qualifiers
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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AR111950
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 25 03-OCT-2000;
                                                                                                                                                                                                                                                     Sequence 25 from patent US
ARI11951
ARI11951.1 GI:12828799
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 24 03-OCT-2000;
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 23 03-OCT-2000;
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Similarity 88.9%;
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                                                                                                          /organism="unknown"
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AR111953/c
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AR111952/c
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Sequence 27
AR111953
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1 (bases 1 to 19)
1 (cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru. 2. O-aminooxy-modified oligonucleotides Patent: US 6127533-A 31 03-OCT-2000;
Location/Qualifiers
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16; Conservat
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Sequence 31 from patent US 6127533.
AR111957
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 27 03-OCT-2000,
Location/Qualifiers
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Cook, P.Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 26 03-OCT-2000;
Location/Qualifiers
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Sequence 26 from patent US
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/mol_type="unassigned
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/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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AR111960/c
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AR111959/c
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AR111959
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Cook, P.Dan., Mancharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 34 03-OCT-2000;
Location/Qualifiers
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 33 03-OCT-2000;
Location/Qualifiers
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Sequence 21
AR124844
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Sequence 20
AR124843
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16; Conserv
l Similarity
16; Conserv
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1 (bases 1 to 19)
Cook, P.Dan., Manoharan, M. and Kawasaki, A. Mamoru.
C'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 44 03-OCT-2000;
                                                                                                                                                Unknown.
Unknown.
                                                                                        Unclassified.

1 (bases 1 to 19)

Manoharan, M., Cook, P.Dan., Prakash, T.P. and Kaw Aminooxy-modified oligonucleotides and methods Patent: US 6172209-A 21 09-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 19)
Manoharan,M., Cook,P.Dan., Prakash,T.P. and Kawasaki,A.M.
Aminooxy-modified oligonuclectides and methods for making
Patent: US 6172209-A 20 09-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unknown.
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larity 88.9%;
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                                           /organism="unknown"
/mol_type="unassigned DNA"
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88.9%;
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Pred. No. 1.9e
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Score 14.8; D
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.ches 2;
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                       DB 1;
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                                                                                                                and Kawasaki, A.
                       Length 19;
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AR124845/c
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AR124847/c
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Manoharan, M., Cook
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Similarity 88.9%; Pred. No. 1.9e+02;
16; Conservative 0; Mismatches 2
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Similarity 88.9%;
  (bases 1 to 19)
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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1 (bases 1 to 19)
Manoharan,M., Cook,P.Dan., Prakash,T.P. and Kawasaki,A.M.
Aminooxy-modified oligonucleotides and methods for making
Patent: US 6172209-A 23 09-JAN-2001;
Location/Qualifiers
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AR124848/c
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AR124849/c
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Best Local Similarity
Matches 16; Conserv
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Sequence 25
AR124848
AR124848.1
                                                                                                                                        1 (bases 1 to 19)
Manoharan, M., Cook, P. Dan., Prakash, T. P. and Kawasaki, A. M. Aminooxy-modified oligonucleotides and methods for making Patent: US 6172209-A 26 09-JAN-2001;
Location/Qualifiers
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Sequence 26 from patent US
AR124849
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Manoharan,M., Cook, P. Dan., Prakash, T.P. and Kawasaki,A.M. Aminooxy-modified oligonuclectides and methods for making Patent: US 6172209-A 25 09-JAN-2001;
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                                                                                                                                                                                                                      Unknown
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nilarity 88.9%;
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25 from patent US 6172209.
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                                                                                                /organism="unknown"
/mol_type="unassigned
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Pred. No. 1.9e+0
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Pred. No. 1.9e+02;
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AR124854/c
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AR124856
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Sequence 27
AR124850
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Manoharan,M., Cook,P.Dan., Prakash,T.P. and Kawasaki,A.M.
Aminooxy-modified oligonuclectides and methods for making same
Patent: US 6172209-A 27 09-JAN-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 19)
Manohara, M., Cook, P. Dan., Prakash, T. P.
Aminooxy-modified oligonucleotides and patent: US 6172209-A 31 09-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 31
AR124854
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                                1 (bases 1 to 19)
Manoharan, M., Cook, P.Dan., Prakash, T.P.
Aminooxy-modified oligonucleotides and π
Patent: US 6172209-A 33 09-JAN-2001;
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/organism="unknown"
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02
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RESULT 194
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AR135291 1
Sequence 20 from patent US
AR135291
AR135291.1 GI:14124196
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Sequence 44 from patent US
AR124867
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Manoharan, M., Cook, P. Dan., Prakash, T. P. and Kawasaki, A. M. Aminooxy-modified oligonuclectides and methods for making Patent: US 6172209-A 44 09-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unclassified.

1 (bases 1 to 19)

Manoharan, M., Cook, P. Dan., Prakash, T. P. and Kawasaki, A. M. Aminooxy-modified oligonucleotides and methods for making Patent: US 6172209-A 34 09-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unknown.
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Sequence 34 from patent US
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/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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AR135292
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Cook, P.Dan., Manoharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonucleotide synthetic intermediates
Patent: US 6194598-A 22 27-FEB-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cook,P.Dan., Manoharan,M. and Kawasaki,A.Mamoru.
Aminooxy-modified oligonucleotide synthetic intermediates
Patent: US 6194598-A 21 27-FEB-2001;
Location/Qualifiers
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Aminooxy-modified oligonucleotide synthetic intermediates
Patent: US 6194598-A 20 27-FEB-2001;
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88.9%; Pred. No. 1.9e+02;
tive 0; Mismatches 2;
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AR135295/c
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Sequence 23
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Cook, P.Dan., Manoharan, M. and Kawasaki, A.Mamoru.

Cook, P.Dan., Manoharan, M. and Kawasaki, A.Mamoru.

Aminooxy-modified oligonuclectide synthetic intermediates

Patent: US 6194598-A 24 27-FEB-2001;

Location/Qualifiers
Unclassified.

1 (bases 1 to 19)
Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
                                                                                                                                                                                                                                                                                                                                                                                                           Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24 from patent US AR135295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 19)
Cook, P.Dan., Manoharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonuclectide synthetic intermediates
Patent: US 6194598-A 23 27-PEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unknown.
                                                                                   AR135296.1
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ilarity 88.9%;
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llarity 88.9%;
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/mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                       /organism="unknown"
/mol_type="unassigned DNA"
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Pred. No. 1.9e
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1.9e+02;
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1.9e+02;
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                    1520 АДАДАДАДАДАДТАДАД 1537
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AR135298
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Sequence
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                                                                                                                                      1 (bases 1 to 19)
Cook, P.Dan., Mancharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonuclectide synthetic inte
Patent: US 6194598-A 27 27-FEB-2001;
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                                                                                      /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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RESULT 202

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AR135305/c
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AR135304/c
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Cook, P.Dan., Manoharan, M. and Kawasaki, A. Mamoru.

Cook, P.Dan., Manoharan, M. and Kawasaki, A. Mamoru.

Aminooxy-modified oligonucleotide synthetic intermediates

Patent: US 6194598-A 31 27-FEB-2001;

Location/Qualifiers
                                1 (bases 1 to 19)
Cook, P.Dan., Manoharan, M. and Kawasaki, A.Mamoru.
Aminooxy-modified oligonuclectide synthetic intermediates
Patent: US 6194598-A 34 27-FEB-2001;
Location/Qualifiers
                                                                                                                                         AR135305 1
Sequence 34 from patent US
AR135305
AR135305.1 GI:14124210
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonucleotide synthetic intermediates
Patent: US 6194598-A 33 27-FEB-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                   Unknown
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Sequence 33 from patent US
AR135304
AR135304.1 GI:14124209
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AR135302
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ilarity 88.9%;
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
             organism="unknown"
/mol_type="unassigned DNA"
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AR141898/c
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AR135315/c
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1520 AAAAAAAAAAAAGTAAAA 1537
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AR153863
Sequence 16 :
AR153863
AR153863.1 (
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AR141898
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AR135315
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Manoharan, M., Kawasaki, A.M., Cook, P.Dan., Fraser, A.S.
Prakash, T.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (bases 1 to 19)
Cook, P. Dan., Mancharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonucleotide synthetic inte
Patent: US 6194598-A 44.27-FEB-2001;
                                                                                                                                                                                                                                            2'-O-acetamido modified monomers and oligomers Patent: US 6147200-A 4 14-NOV-2000;
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/mol_type="unassions
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/mol_type="unassigned
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
0; Mismatches 2
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AR164173/c
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Best Local Similarity
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 GENSET
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PD 1:
PF 2:
PR 2:
                                                                               Cohen,D., Blumenfeld,M., Chumakov,I. and Bougueleret,L. Prostatic cancer gene Patent: JP 2002516657-A 489 11-JUN-2002;
                                                                                                                     Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 19)
                                                                                                                                                                         BD196900.1 GI:33006670
JP 2002516657-A/489.
Homo Bapiens (human)
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Prostatic cancer gene.
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Manoharan,M., Mohan,V. and Boswell,H.
RNA targeted 2'-modified oligonucleotides that are conformationally
preorganized
Patent: US 6271358-A 6 07-AUG-2001;
Location/Qualifiers
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Sequence 6 from
AR164173
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Heller, M.J., Tu, E., Evans, G.A. and Sosnowski, R.G.
Methods for transport in molecular biological analysis
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Homo sapiens (human)

DP 2002516657-A/489

11-JUN-2002

12-DEC-1998 JP 2000525562

22-DEC-1997 US 08/996306,09-SEP-1998
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Pred. No. 1.9e+02;
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PC C12N5/00,C12N15/00
PC C12N5/00,C12N15/00
PC potential microsequencing oligo for 4-4-187.mis2 FH Key Location/Qualifiers
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PN JP 2002516657-A/500

PD 11-JUR-2002

PD 22-DEC-1998 UP 2000525562

PF 22-DEC-1997 US 08/996306,09-SEP-1998 US 60/099658 P

DANIEL COHEN, MARTA BLUMENFELD, ILYA CHUMAKOV, LYDIE BOUGUELERET

C12N15/09, C12N15/09, A01K67/027, C07K14/47, C07K16/18, C12N1/15, P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cohen,D., Blumenfeld,M., Chumakov,I. and Prostatic cancer gene Patent: JP 2002516657-A 500 11-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 19)
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PC C12N5/00,C12N15/00
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Similarity 88.9%;
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                                                                                                                            /organism="Homo sapiens"
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/db_xref="taxon:9606"
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RESULT 211 BD274438/c

LOCUS DEFINITION

BD274438 19 l Oligonucleotides having A-DNA

A form

DNA and

linear PAT 17-JUL-2 B-DNA form confirmational

PAT 17-JUL-2003

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REFERENCE
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BD274439/c
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Oligonucleotides having A-DNA
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synthetic construct
artificial sequences.

1 (bases 1 to 19)
Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA form
                                                                                                                                                                            synthetic construct
synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 19)
Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA form confirmational
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ISIS PHARMACEUTICALS INC
OS Artificial Sequence
PN JP 2002543215-A/15
                                                                                                                                                                       geometry
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 C12N15/00
Oligonucleotide
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2' - O-MOE linkage
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misc_feature
misc_feature
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03-MAY-2000 JP 2000615638
03-MAY-1999 US 09/303586
MUTHIAH MANCHARAN, VENKATRAMAN MOHAN
C07H21/02, A61K48/00, A61P35/00, A61P35/02, A61P43/00, C12N15/09,
                                                                                                         nt: JP 2002543215-A 16 17-DEC-2002;
PHARMACEUTICALS INC
Artificial Sequence
JP 2002543215-A/16
17-DEC-2002
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llarity 88.9%;
Conservative
                                                           03-MAY-2000 JP 2000615638
03-MAY-1999 US 09/303586
MUTHIAH MANOHARAN, VENKATRAMAN MOHAN
COTH21/02, A61K48/00, A61P35/00, A61P35/02, A61P43/00, C12N15/09,
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/db_xref="taxon:32630"
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(17) . . (18)
(18) . . (19) .
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Pred. No. 1.9e
0; Mismatches
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and
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B-DNA form confirmational
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 19;
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   RESULT 214
BD274441/c
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REFERENCE
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VERSION
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BD274440/c
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                                             Matches
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Best Local
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Best Local :
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19
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                                             16;
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1 (bases 1 to 19)
Manoharan,M. and Mohan,V.
Oligonucleotides having P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BD274440
Oligonucleotides having A-DNA
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                                                                                                                                                                                                                                                                                                                                                                                                        Patent: JP 2002543215-A 17 17-DEC-2002;
                                                         Similarity
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AAAAAAAAAAAAAAAAA
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Artificial Sequence
JP 2002543215-A/17
17-DEC-2002
                                                                                                                                                     misc_feature
misc_feature
misc_feature
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03-MAY-1999 US 09/301586
MUTHIAH MANOHARAN, VENKATRAMAN MOHAN
COTH21/02,A61K48/00,A61P35/00,A61P35/02,A61P43/00,C12N15/09,
C12N15/00
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misc_feature (17)...
misc_feature (18)...
Location/Qualifiers
                                             Conservative
                                                                                                                                                                                     misc_feature
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3' - O-MOE linkage; sub O linkage
3' - O-MOE linkage; sub O linkage
3' - O-MOE linkage
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sub O linkage
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                                                                                        /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                        Location/Qualifiers
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/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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Pred. No. 1.9e-
0; Mismatches
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                                            Score 14.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2;
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.9e+02;
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BD274449/c
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OS Artificial Sequence
PN JP 2002543215-A/26
PD 17-DEC-2002 JP 200061
PF 03-MAY-1909 US 09/
PI MUTHIAH MANOHARAN, VEN
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Patent: ,
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Patent: (
                                                                                                                                                                           synthetic construct
synthetic construct
artificial sequences.

1 (bases 1 to 19)
Manoharan, M. and Mohan, V.
Oligonucleotides having A-DNA form and B-DNA form confirmational
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Oligonucleotides having
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synthetic construct
artificial sequences.
1 (bases 1 to 19)
Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA
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JP 2002543215-A/18.
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Oligonucleotides having A-DNA
                                                                                                                                                                                                                                                                                                           JP 2002543215-A/26.
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Similarity 88.9%;
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misc_feature
misc_feature
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MUTHIAH MANOHARAN VENKATRAMAN MOHAN
C07H21/02,A61K48/00,A61P35/00,A61P35/02,A61P43/00,C12N15/09,
C12N15/00
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03-MAY-2000 JP 2000615638
03-MAY-1999 US 09/3035
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PHARMACEUTICALS INC
Artificial Sequence
JP 2002543215-A/18
  17-DEC-2002
03-MAY-2000 JP 2000615638
03-MAY-1999 US 09/303586
03-MAY-1999 US VENKATRAMAN MOHAN
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misc_feature
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2'- O-MOE; sub O linkage
2'- O-MOE; sub O linkage
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                                                                                                                                            JP 2002543215-A 26 17-DEC-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No. 1.9e+02;
0; Mismatches 2;
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and B-DNA form
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VERSION
KEYWORDS
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VERSION
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AR205799/c
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AR205798/c
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1520 AAAAAAAAAAAAGTAAAA 1537
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Sequence 16
AR205799
AR205799.1
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16; Conservative
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AR205798
geometry
Patent: US 6369209-A 16 09-APR-2002;
Location/Qualifiers
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Patent: US 6369209-A 15 09-APR-2002;
Location/Qualifiers
                                               1 (bases 1 to 19)
Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and
                                                                                                           Unknown
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Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and
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2'-modified T linkage
2'-modified T linkage
2'-modified T linkage
2'-modified T linkage
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                                                                                                                                                                                                                                                                                                                                                                /mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                                                                                                                                                                                                                                                                                                                                                                 organism="unknown"
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Pred. No. 1.9e+02;
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Pred. No. 1
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                                                     B-DNA form conformational
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RESULT 220
AR205809/c
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AR205800/c
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AR205801/c
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Patent: L
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AR205801
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
AR205800
  AR205809
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Manoharan, M. and Mohan, V.
Oligonucleotides having A-DNA form and B-DNA
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Similarity 88.9%;
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 1.9e.
0; Mismatches
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Pred. No. 1.9e+02;
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AR213490/c
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1520 ААЛАЛАЛАЛАЛАГТАЛАЛ 1537
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Sequence 1 from p
AR213490
AR213490.1 GI:23
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Sequence 2
AR213491
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Patent: US 6403779-A 1 11-JUN-2002;
Location/Qualifiers
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Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA form conformational
                          Regioselective synthesis of 2'-O-modified nucleosides
Patent: US 6403779-A 2 11-JUN-2002;
Location/Qualifiers
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AR205809
                                                                                     1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
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Rawasaki,A.M., Fraser,A.S., Manoharan,M., Cook,P.D. and
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                                                                              Prakash, T.P.
                                                                                                                            Unclassified
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/mol_type="genomic DNA"
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/mol_type="unassigned DNA"
/organism="unknown"
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Pred. No. 1.9e+02;
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AR213494/c
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AR213493/c
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AR213494
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Sequence 4 from patent US
AR213493.
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                                                                                                                                                                                                                                 Unclassified.
1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
                                                                                                                                                                                                                                                                                Unknown
                                                                                                                                                                                                                                                                                                                  AR213493.1 GI:23310724
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1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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llarity 88.9%;
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/mol_type="genomic DNA"
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AR213496/c
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Unclassified.
                                               Regioselective synthesis of 2'-O-modified Patent: US 6403779-A 7 11-JUN-2002;
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Kawasaki, A.M., Fraser, A.S., Manoharan, M., Cook, P.D. and
Prakash, T.P.
                                                                                                                         Unknown
                                                                                                                                   Unknown.
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Kawasaki, A.M., Fraser, A.S., Manoharan, M., Cook, P.D. and Prakash, T.P.
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Similarity 88.9%; Pred. No. 1.99
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AR213497/c
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AR213502/c
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AR213501/c
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AR213502
Sequence 14 from patent US 6403779.
AR213502
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Sequence Brom patent US 6403779.
AR213497 GI:23310728
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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AR213512
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Kawasaki A.M., Fraser, A.S., Manoharan, M.,
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Kawasaki, A.M., Fraser, A.S., Manoharan, M.,
Prakash, T.P.
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Pred. No. 1.9e+02;
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                                                                                        Cook, P.D. and
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                                          RESULT 235
AR295557/c
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KEYWORDS
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AR237463/c
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Kurz,M., Lohse,P. and Wagner,R.

Peptide acceptor ligation methods

Patent: US 6429300-A 25 06-AUG-2002;

Location/Qualifiers
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Sequence
AR222465
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16; Conserv
                    AR295557
Sequence 7292 from patent
AR295557
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Sequence 1 from patent US
AR237463
AR237463.1 GI:27282213
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Ravikumar, V.T., Manoharan, M., Capaldi, D.C., Krotz, A.,
Guzaev, A.
                                                                                                                                                                                                                                                                                                                                                                                                                            1 AAAAAAAAAAAAAAAAA 18
          AR295557.1 GI:31682841
                                                                                                                                                                                                                     Process for the synthesis of oligomeric Patent: US 6465628-A 1 15-OCT-2002; Location/Qualifiers
                                                                                                                                                                                                                                                                                       Unclassified.
                                                                                                                                                                                                                                                                                                    Unknown
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ilarity 88.9%;
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ilarity 88.9%;
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/mol_type="genomic DNA"
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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AR359804/c
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AR321589/c
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                                                                                          1 (bases 1 to 19)
Manoharan,M. Cook,P.D., Prakash,T.P. and Mohan,V.
Guanidinium functionalized nucleotides and precurs
Patent: US 6593466-A 3 15-JUL-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
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Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing a high density
disequilibrium map of the human genome
Patent: US 6537751-A 7292 25-MAR-2003;
                                                                                                                                                                             Unknown
                                                                                                                                                                                                                                  Sequence 3 from patent US AR359804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 19)
Baxter,A.D., Collingwood,S.P., Douglas,M.E. and Taylor,R.J.
Oligonucleotide analogues
Patent: US 6562960-A 10 13-MAY-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AR321589
Sequence 10 from patent US
AR321589
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Similarity 88.9%;
16; Conservative
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Similarity 88.9%;
16; Conservative
            Similarity
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                                                    /organism="unknown"
/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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1.1%; Score 14.8; DB 1;
88.9%; Pred. No. 1.9e+02;
vative 0; Mismatches 2;
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Pred. No. 1.9e+02;
0; Mismatches 2
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Pred. No. 1.9e+02;
0; Mismatches 2;
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AR359805/c
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AR367447/c
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AR359806/c
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Matches 16; Conserv
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Sequence 4 f
AR367447
AR367447.1
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Sequence
AR359805
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16; Conserv
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Manoharan,M., Cook,P.D., Prakash,T.P. and Mohan,V.
Guanidinium functionalized nucleotides and precurs
Patent: US 6593466-A 4 15-JUL-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                         1 (bases 1 to 19)
Mancharan,M., Cook,P.D., Prakash,T.P. and Mohan,V.
Guanidinium functionalized nucleotides and precurs
Patent: US 6593466-A 5 15-JUL-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                               AR359806
Sequence 5 from patent 1
AR359806
AR359806.1 GI:33766604
1 (bases 1 to 19)
Collingwood, S.P., Moser, H.E.,
                                                                                                                                                                                                                                                                                                                                                              Unknown
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                                                                                             4 from patent US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mol_type="genomic DNA"
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                                                                                                                                                                                                                                                                      organism="unknown"
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                                                                         GI:34600659
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                                                                                                                                                                                                             Score 14.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2
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Pred. No. 1.9e+02;
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    Altmann, K.-H.
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AR399177/c
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AR399178/c
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AR399178
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AR399177
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Crooke, S.T., Lima, W.F., Wu, H. and Monoharan, M.
Human Rnase H1 and oligonucleotide composition
Patent: US 6617442-A 18 09-SEP-2003;
                                                                                                                                                                                                        Unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 19)
Crooke, S.T., Lima, W.F., Wu, H. and Monoharan, M. Human Rnase H1 and oligonucleotide composition Patent: US 6617442-A 17 09-SEP-2003;
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Similarity 88.9%; Pred. No. 1.9¢
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/mol_type="genomic |
                                                                                          /organism="unknown"
/mol_type="genomic |
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/mol_type="genomic DNA"
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88.9%; Pred. No. 1
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88.9%; Pred. No. 1.90
tive 0; Mismatches
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RESULT 243

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AR403602/c
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AR403603/c
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AR403602
AR403602.1 GI:40151188
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AR403601
AR403601.1 GI:40151187
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Sequence 3 from patent US
AR403603
AR403603.1 GI:40151189
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Regioselective synthesis of 2'-O-modified Patent: US 6624294-A 3 23-SEP-2003; Location/Qualifiers
                                 1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
                                                                                       Unknown.
                                                                                                                                                                                                                                                                                                                                    Regioselective synthesis of 2'-O-modified Patent: US 6624294-A 2 23-SEP-2003;
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1 (bases 1 to 19)
Kawasaki.A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Draitach T
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ilarity 88.9%;
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/mol_type="genomic DNA"
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AR403604/c
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AR403606/c
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Sequence 5
AR403605
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unclassified.

(bases 1 to 19)

(bases 1 to 19)

(Kawasaki, A.M., Fraser, A.S., Manoharan, M., Cook, P.D. ar Prakash, T.P.

Regioselective synthesis of 2'-O-modified nucleosides Patent: US 6624294-A 23-SEP-2003;
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AR403604
AR403604.1 GI:40151190
                                                                                                                                                                                          Unclassified.
1 (bases 1 to 19)
Kawasaki, A.M., Fraser, A.S., Manoharan, M.,
Prakash, T.P.
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Local Similarity 88.9%;
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Sequence 6
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AR403608
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        Regioselective synthesis of 2'-O-modified nucleosides
Patent: US 6624294-A 8 23-SEP-2003;
Location/Qualifiers
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
                                          Unclassified.
1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M., Cook,P.D. and Prakash,T.P.
                                                                                     Unknown
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Kawasaki, A.M., Fraser, A.S., Manoharan, M.,
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Unclassified.
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/mol_type="genomic DNA"
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Kawasaki, A.M., Fraser, A.S., Manoharan, M.,
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1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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Similarity 88.9%;
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/mol_type="genomic |
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/mol_type≂"genomic DNA"
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AR412338/c
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AR403623/c
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Cook, P.D., Manoharan, M., Maier, M. and An, H.
C3'-methylene hydrogen phosphonate oligomers and
Patent: US 6639061-A 1 28-OCT-2003;
Location/Qualifiers
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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1 (bases 1 to 19)

Kawasaki, A.M., Fraser, A.S., Mancharan, M., Prakash, T.P.
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/organism="unknown"
/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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         AR451262
Sequence 5 !
AR451262
AR451262.1
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Sequence 7
AR432617
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1 (bases 1 to 19)

Manoharan,M., Cook,P.D. and Guinosso Modified oligonucleotides
Patent: US 6653458-A 6 25-NOV-2003;
Location/Qualifiers
                                                                                                                                                                                                                  Unclassified.
1 (bases 1 to 1)
Manoharan, M., Cook, P.D. and Guinosso
Modified oligonucleotides
Patent: US 6653458-A 7 25-NOV-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unknown
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AR432616
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Similarity 88.9%;
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                                                                                                                                                                        /organism="unknown"
/mol_type="genomic DNA"
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/mol_type="genomic"
                                                                                                                                                                                                        Location/Qualifiers
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           GI:42682240
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Pred. No. 1.9e+02;
0; Mismatches 2;
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Pred. No. 1.9e+02;
0; Mismatches 2
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Pred. No. 1.9e+02;
                                                                                                                                                                                                                                        and Guinosso, C.
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AX059378/c
LOCUS
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KEYWORDS
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AR451282/c
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Best Local Similarity
Matches 16; Conserv
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Best Local
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Query Match
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Sequence 26 from patent US 6673912.
AR451282
AR451282.1 GI:42682260
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Mancharan,M. and Cook,P.D.
2'-O-aminoethyloxyethyl-modified oligonucleotides
Patent: US 6673912-A 5 06-DAN-2004;
Location/Qualifiers
                                                                                                   Plant chromosome compositions and methods Patent: WO 0055325-A 111 21-SEP-2000; The University of Chicago (US)
                                                                                                                                                                        Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
Arabidopsis thaliana
Eukaryotopis thaliana
Eukaryotopis Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Sequence 111 from Patent WO0055325.
AX059378
AX059378.1 GI:12311483
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Manoharan, M. and Cook, P.D.
2'-O-aminoethyloxyethyl-modified oligonucleotides
Patent: US 6673912-A 26 06-JAN-2004;
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/mol_type="genomic DNA"
                              /organism="Arabidopsis thaliana"
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/db_xref="taxon:3702"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="unknown"
/mol_type="genomic DNA"
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O; Mismatches
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Pred. No. 1.9e+02;
 Score 14.8;
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AX132398/c
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DEFINITION
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AX226133
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16; Conservative
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                                                                                                                                                                                                                                                                   synthetic construct
synthetic construct
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Ribozyme therapy for the treatment of proliferative skin and
                                                                                                                                                               Patent: WO 0160856-A 52 23-AUG-2001;
UNIVERSITE CATHOLIQUE DE LOUVAIN (BE)
Location/Qualifiers
                                                                                                                                                                                                                 vmglom gene and its mutations causing disorders
                                                                                                                                                                                                                                                     artificial sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent: WO 0130362-A 3616 03-MAY-2001; IMMUSOL, INC. (US)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Similarity 88.9%;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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                                                                                                 /organism="synthetic construct"
/mol_type="unassigned DNA"
/mol_tyse="taxon:32630"
/mole="oligonucleotide"
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                                                           1.1%;
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                                               Score 14.8; D. Pred. No. 1.9e. 0; Mismatches
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Pred. No. 1.9e+02;
0; Mismatches
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D; Mismatches 2;
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1.9e+02;
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AX349249/c
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                                                              FEATURES
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Matches 16; Conserv
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CC with dyes
FH Key L
FT source 1
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                                                                                                                                                                                                                                                                 Sali-assembling microelectronic integration system capable of designating self address, compartment device, mechanism, metho operation for molecular biological analysis and diagnosis Patent: UP 2001525193-A 16 11-DEC-2001;

NANOGEN INC
OS Artificial Sequence
PN UP 2001525193-A/16
PD 11-DEC-2001
PP 01-DEC-2001
PP 01-DEC-2001
PR 05-DEC-1998 UP 2000524303
PR 05-DEC-1998 UP 200524303
PR 05-DEC-1998 UP 200524303
PR NOALD G SOSNOWSKI, WILLIAM F BUTLER, EUGENE TU, MICHAEL I
                                                                                                                                                                                        NERENBERG,
PI MICHA
PC C12Q1
CC Descr
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BD087505
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Schulz,T. and Wagner,G.
Method for qualitative and/or quantitative detecting of molecular interactions on probe arrays
Patent: WO 0202810-A 33 10-JAN-2002;
Clondiag Chip Technologies GmbH (DE)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                artificial sequences.
1 (bases 1 to 19)
Sosnowski,R.G., Butler,W.F., Tu,E., Nerenberg,M.I., Heller,M.J.
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synthetic construct
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AX349249
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Description of Artificial Sequence:
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llarity 88.9%;
Conservative
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                                                      /organism='Artificial Sequence'
/organism="synthetic construct"
/mol_type="genomic DNA"
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/mol_type="unassigned DNA"
/db xrefe"taxon:32630"
/noTe="Oligonukleotidsonde"
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from Patent WO0202810.
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Pred. No. 1.9e+02;
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RESULT 267
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AR064875/c
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1 (bases 1 to 20)
Holton, T. Albert., C
Lester, D. Ruth.
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AR080000 20 bp
Sequence 83 from patent US 5968524.
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                                                                                                                                                                                                                                                                      Unknown
                                                                                                                                                                                                            1 (bases 1 to 20)
Cros, P., Kurfurst, R., Battail, N. and Piga, N.
Process and device for assaying a hapten
Patent: US 5849480-A 5 15-DEC-1998;
                                                                                                                                                                                                                                                                                                                    Sequence 5
AR064875
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Location/Qualifiers
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AR030917
                                                                                                                                                                                                                                                                                 Unknown.
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                                                                                                                                                           /organism="unknown"
/mol_type="unassigned DNA"
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88.9%; Pred. No. 1.9e+02;
vative 0; Mismatches 2
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                                                                                                                            Score 14.8; DB 1;
Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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Mismatches 2
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             31-AUG-2000
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AR087520/c
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Sequence 1
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Sequence 83
AR085926
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                                     i (bases 1 to 20)
Pitsch,S., Weiss,P.A. and Jenny,L.
Ribonucleoside-derivative and method for preparing the
Patent: US 5986084-A 1 16-NOV-1999;
Location/Qualifiers
                                                                                                Unknown
                                                                                                                                                                                                                                                                                                                          Patent: US 5985287-A 83 16-NOV-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 20)
Tan, P., Skinner, M. and
Compounds and methods
                                                                                                                                                                                                                                                                                                                                                                                              Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent: US 5968524-A 83 19-OCT-1999;
Location/Qualifiers
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1 (bases 1 to 20)

Watson, J.D. and Tan, P.
Methods and compounds
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AR080000.1
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                                                                                                                                                                                                                                                                                          /mol_type="unassigned
          /mol_type="unassigned
                   organism="unknown"
                                                                                                                                                                                                                                                                                                      organism="unknown"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="unknown"
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Pred. No. 1.8e+02;
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          AR121540
Sequence
AR121540
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AR118970
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AR093312
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Tan, P., Hiyama, J., Visser, E., Skinner, M.,
Mycobacterium vaccae antigens
Patent: US 6001361-A 83 14-DEC-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unknown
                                                                                                                                                                                                                       1 (bases 1 to 20)
Uchida, K., Uchida, T., Tanaka, Y., Matsuda, Y.
Antisense nucleic acid compound targeted to
Patent: US 6150092-A 96 21-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                      1 AAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                              Unknown
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AR121540.1
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Similarity 88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="unknown"
/mol_type="unassigned
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/mol_type="unassigned
                                                                                                                                                                                                            Location/Qualifiers
                    from
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Pred. No. 1.8e
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Pred. No. 1.8e+02;
0; Mismatches 2;
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AR123335
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AR121692
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Query Match
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Matches 16
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TITLE
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Best Local Similarity
Matches 16; Conserv
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Sequence 83
AR121692
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MCKAY, R., Borchers, A.H. and Baker, B.F.

Antisense modulation of peroxisome proliferator-activated receptor gamma expression

Patent: US 6159734-A 76 12-DEC-2000;

Location/Qualifiers
                                                                         1 (bases 1 to 20)
Bruice, T.C. and Dev, A.P.
Deoxynucleic alkyl thiourea compounds and
Patent: US 6169176-A 1 02-JAN-2001;
Location/Qualifiers
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1.1%; Score 14.8; DB 1; Similarity 88.9%; Pred. No. 1.8e+02; 16; Conservative 0; Mismatches 2;
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Sequence 1 from patent US
AR123335
                                                                                                                                                                                                                                                                                                                                                                                                       Patent: US 6160093-A 83 12-DEC-2000; Location/Qualifiers
                                                                                                                                                 Unknown
                                                                                                                                                                                    AR123335.1
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                                                                                                                                       Unclassified.
                                                                                                                                                            Unknown.
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/mol_type="unassigned
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/mol_type="unassigned DNA"
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Pred. No. 1.8e+02;
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AR139962/c
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AR139960/c
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                                                                                                                RESULT 277
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Query Match 1.1%;
Best Local Similarity 88.9%;
Matches 16; Conservative
  AUTHORS
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                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 20)
Zsebo, K.M., Bosselman,R.A., Suggs,S.V.
DNA encoding stem cell factor
Patent: US 6207417-A 34 27-MAR-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 34 from patent US AR139962
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AR139960
Unclassified.
1 (bases 1 to 20)
Zsebo,K.M., Bosselman,R.A.,
                                            Unknown.
                                                                                                 AR140279
Sequence 32 from patent
AR140279
                                                                                                                                                                                                                                                                                                                                                                                                                                              Unknown
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Zsebo, K.M., Bosselman, R.A., Suggs, S.V.
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ilarity 88.9%;
Conservative
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/mol_type="unassigned DNA"
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; Pred. No. 1.8e
0; Mismatches
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Pred. No. 1.8e+02;
0; Mismatches 2
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s 6207454
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S 6207417.
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 Suggs, S.V. and Martin, F.H.
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AR140557/c
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AR140281/c
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Sequence 32 from patent US
AR140557
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Seebo, K.M., Bosselman, R.A., Suggs, S.V. and Martin, F.H.

Method for enhancing the effciency of gene transfer with stem
factor (SCF) polypeptide
Patent: US 6207454-A 34 27-MAR-2001;
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                                                                                                                                                       1 (bases 1 to 20)
25 (bases 1 to 20)
26 (bo, K.M., Bosselman, R.A., Suggs, S.V. Stem cell factor and compositions Patent: US 6207802-A 32 27-MAR-2001;
                                                                                                                                                                                                                       Unknown
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Similarity 88.9%;
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                                                                                                   /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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1.8e+02;
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AR140559/c
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AR154115/c
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AR141070/c
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AR154115
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AR140559
Huang, Z. and Szostak, J.W.

Simple and efficient method to label and modify 3'-termini of RNA using DNA polymerase and a synthetic template with defined overhan nucleotides
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Manoharan, M. and Maler, M.A.
Compounds, processes and intermediates
backbone oligomeric compounds
Patent: US 6207819-A 1 27-MAR-2001;
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Stembo, K.M., Bosselman, R.A., Suggs, S.V.
Stem cell factor and compositions
Patent: US 6207802-A 34 27-MAR-2001;
Location/Qualifiers
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1 (bases 1 to 20)
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Best Local Similarity 88.9%;
Matches 16; Conservative
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PALE PATENT: JP 2002514385-A 26 21-MAY-2002;
GENESIS RESEARCH AND DEVELOPMENT CORP LTD

OS Artificial Sequence
PN JP 2002514385-A/26
PD 21-MAY-2002
PF 23-DEC-1998 JP 2000525553
PR 23-DEC-1997 US 08/997362,23-DEC-1997 US 08/997080 PR
23-DEC-1997 US 08/996624,11-JUN-1998 US 09/095855 PR
23-DEC-1997 US 08/996624,11-JUN-1998 US 09/205426 PI PAUL PRESP-1998 US 09/1956181,04-DEC-1998 US 09/205426 PI PAUL PRESTICCE
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PC C12N15/09,A61K31/711,A61K39/04,A61K48/00,A61P11/00,A61P11/00, A61P17/00,
PC A61P17/00,
PC A61P17/00,
PC C7K19/00,
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Sequence 13
AR164658
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Compositions derived from
their use.
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1 (bases 1 to 20)

Tan,P., Watson,J., Visser,E.S., Skinner,M.A. and Prestid,R.L.

Compositions derived from mycobacterium vaccae and methods for
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Blumberg, B.
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Location/Qualifiers
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     A61P17/00,
A61P17/06, A61P31/00, A61P31/06, A61P37/04, C07K14/35, C07K16/12,
C07K19/00,
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Protein skeleton of antibody
BD234126
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 synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  artificial sequences.
1 (bases 1 to 20)
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Similarity 88.9%;
16; Conservative
                                                                                                                                 AAAAAAAAAAAAAAAA 18
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C12N15/00
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                                                                                                                                                                                                                                                                                                                                                                                                             Artificial Sequence
JP 2002532072-A/14
02-0CT-2002
09-DEC-1999 JP 2000587187
10-DEC-1998 US 60/111737
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Key
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                                                                                                                                                                                                                                                                                                                                Puromycin linker oligonucleotide
Key Location/Qualifiers
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                                                                                                                                                                                                                                    /organism="synthetic co
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                         /organism='Artificial Sequence'
Location/Qualifiers
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                           GI:44849560
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                                                    from Patent
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Pred. No. 1.8e+02;
0; Mismatches 2;
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Pred. No. 1.8e+02;
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WO2003106672.
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JUNICLEOTIDE

SOVANCY GIUTSU KENKYUSHO:KK

ANTI-HTLV-I ANTI-SENSE OLIGONUCLEOTIDE

PAtent: JP 1997052898-A 10 25-FEB-1997;

SOVANCY GIUTSU KENKYUSHO:KK

OS None

PH UP 1997052898-A/10

PD 25-FEB-1997

PF 09-AUG-1995 JP 1995224606

PI MIZUGUCHI MASATSUGU, KUROSAKI NAOKO

KOYANGI YOSHIO,

PC CO7H21/04/ASIK31/70;

CC topology: Linear;

CC topology: Linear;

CC anti-sense:

PH Key

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Patent: WO 2003106672-A 40 24-DEC-2003;
Riken (JP) ; Kabushiki Kaisha Dnaform (JP)
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                                                                                                                                            /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol type="unassigned DNA"
/db xref="taxon:32630"
/noTe="tag8"
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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Sequence 11 from patent US 5639870.
147310
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Sequence 16 from patent US
I36180
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Holton,T.A., Cornish,E.C., Kovacic,F., Tanaka,Y. and Lester,D.R.
Genetic sequences encoding flavonoid pathway enzymes and uses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 from pate
128309
128309.1 GI:1819085
                                           Holton,T.Albert., Cornish,E.Cecily. and Tanaka,Y. Genetic sequences encoding flavonoid pathway enzymes
                       Patent: US 5639870-A 11 17-JUN-1997;
                                                         1 (bases 1 to 20)
Holton, T. Albert.,
                                                                                            Unknown
                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 20)
Heller, M.J. and Tu, E.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Unknown
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/mol_type="unassigned
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/mol_type="unassigned
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Pred. No. 1.8e+02;
0; Mismatches 2;
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AR222466
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AR231312
Sequence
                                                                                                                                                                                              Unclassified.
1 (bases 1 to 20)
Kurz,M., Lohse,P. and Wagner,R.
Peptide acceptor ligation methods
Patent: US 6429300-A 26 06-AUG-2002;
                                                                                                            l Similarity
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Sequence 26 from patent US
AR222466
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1 (bases 1 to 20)

1 (bases 1 to 20)

1an, P., Visser, E., Prestidge, R. and Watson, J.D.

Compounds and methods for treatment and diagnosis of mycobacterial
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Sequence 83 from patent US
AR213738
                                                                                                                                                                                                                                                       Unknown.
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llarity 88.9%;
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larity 88.9%;
Conservative C
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49 from
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/mol_type="genomic DNA"
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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Egholm,M., Nielsen,P., Buchardt,O.,
Coull,J.M., Kiely,J. and Griffith,M.
Peptide nucleic acids
Patent: US 6451968-A 49 17-SEP-2002;
                                                                1 (bases 1 to 20)
Mirkin,C.A., Letsinger,R.L., Mucic,R.C.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides att
                                                                                                                                                                                                     Sequence 55
AR274394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compounds, processes and intermediates backbone oligomeric compounds Patent: US 6462184-A 1 08-OCT-2002; Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unknown.
                                                                                                                                                 Unknown.
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Manoharan, M. and Maier, M.A.
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AR236083
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                            Patent: US 6506564-A 55 14-JAN-2003;
Location/Qualifiers
                                                                                                                                Unclassified.
                                                                                                                                                             Unknown.
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Similarity 88.9%;
16; Conservative
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/mol_type="genomic DNA"
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/mol_type="genomic |
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/organism="unknown"
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84.2%; Pred. No. 1.8e+02;
tive 0; Mismatches 3
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Pred. No. 1.8e+02
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                                                                          attached thereto and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for
                                                                                                      Storhoff, J.J.,
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RESULT 298
AR365970
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                                                                                                                                   Ch 1.1%; Score 14.8; DB 1; 1 Similarity 88.9%; Pred. No. 1.8e+02; 16; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                      1 (bases 1 to 20)
Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides attached thereto and uses
thereof
AR365970 20 bp
Sequence 83 from patent US 6328978.
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                                                                                                                                                                                                                            Patent: US 6582921-A 55 24-JUN-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                      Unknown
                                                                                                                                                                                                                                                                                                                                                   Unknown.
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Sequence 55 from patent US
AR344936
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Manoharan, M., Lonnberg, H., Salo, H. and Virta, P. Aminooxy functionalized oligomers
Patent: US 6576752-A 10 10-JUN-2003;
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Unclassified.
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                                                                                ARAAAAAAAAAAAA 18
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                                                                                                                                                                                     /organism="unknown"
/mol_type="genomic |
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AR382312
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Sequence
AR429653
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Unknown.
                                                               1 (bases 1 to 20)
Mirkin, C.A., Letsinger, R.L., Mucic, R.C.,
Elghanian, R. and Taton, T.A.
                                                                                                                                                                                                                                                                         1 ААААААААААААААААА 18
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Mirkin, C.A., Letsinger, R.L., Mucic, R.C., Storhoff, J.J.,
Elghanian, R. and Taton, T.A.
Nanoparticles having oligonucleotides attached thereto and uses
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Watson, J.D., Tan, P.L.J. and Prestidge, R.
Methods for the treatment of immunologically-mediated skin
disorders
                                                                                                                        Unknown
                                                                                                                                                               AR429653.1
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Location/Qualifiers
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AR365970.1
                                         Nanoparticles having oligonucleotides therefor
                                                                                                                                      Unknown.
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Location/Qualifiers
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             Patent: US 6645721-A 55 11-NOV-2003;
Location/Qualifiers
                                                                                                         Unclassified.
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Similarity 88.9%;
16; Conservative
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/mol_type="genomic |
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Pred. No. 1.8e+02;
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5 6610491.
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                                                                                 Storhoff, J.J.,
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Sequence 55
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1 (bases 1 to 20)
Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J., Elghanian,R. and Taton,T.A.
Ranoparticles having oligonucleotides attached thereto
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Sequence 55 from patent US
AR447441
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                                                                                                                                   Patent: US 6677122-A 55 13-JAN-2004;
Location/Qualifiers
                                                                                                                                                                        1 (bases 1 to 20)
Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides attached thereto
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 AAAAAAAAAAAAAAAAA 18
                                              1.1%;
ilarity 88.9%;
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ilarity 88.9%;
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/mol_type="genomic DNA"
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/mol_type="genomic |
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/mol_type="genomic DNA"
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; Pred. No. 1.8e.
0; Mismatches
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; Pred. No. 1.8e-
0; Mismatches
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Pred. No. 1.8e+02;
0; Mismatches 2
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RESULT 303

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Sequence 55
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Sequence 7
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1 (Dases 1 to 20)
1 (Dases 1 to 20)
Mirkin, C.A., Letsinger, R.L., Mucic, R.C., Storhoff, J.J., Elghanian, R. and Taton, T.A.
Nanoparticles having oligonucleotides attached thereto
                                                                  Unknown
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Nelson, W.G., Lin, X., Tchou, J.C. and Bakker, J.
Methods of diagnosing and treating hepatic ce
                                                                                                                                                                                                                                                                                                                                                                                                                           Unknown
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Mirkin, C.A., Letsinger, R.L., Mucic, R.C.,
Elghanian, R. and Taton, T.A.
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Sequence 55 from patent
AR454776
                                                                             Unknown.
                                                                                                                                                                                                                                                                                                                                    Patent: US 6709818-A 7 23-MAR-2004;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                       Unknown.
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/mol_type="genomic DNA"
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1 (bases 1 to 20)
Mirkin, C.A., Letsinger, R.L., Mucic, R.C., Storhoff, J.J., Elghanian, R. and Taton, T.A.
Nanoparticles having oligonucleotides attached thereto
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                                                                                                                                    Patent: US 6720411-A 55 13-APR-2004;
Location/Qualifiers
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Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides attached thereto
                                                                                                                                                                                                                                      Unknown
                                                                                                                                                                                                                                                                                      Sequence 55 from patent US AR494728
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Location/Qualifiers
                                                                                               /mol_type="genomic DNA"
                                                                                                            organism="unknown"
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Pred. No. 1.8e+02;
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AX045779/c
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ACCESSION
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AX004876/c
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AX045787/c
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Sequence 9 from Patent WO0067023.
AX045779
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Method for isolating anionic organic substances
systems using carionic polymer nanoparticles
Patent: WO 9910527-A 5 04-MAR-1999;
SUEDDEUTSCHE KALKSTICKSTOFF (DE); BAYER ERNST (DE)
Location/Qualifiers
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artificial sequences.
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Sequence 5 from Patent WO9910527.
AX004876
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Similarity 88.9%;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Pred. No. 1.8e+02;
0; Mismatches 2;
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AX045790/c
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AX056597/c
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Best Local Similarity
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Murinae gen. sp.
Murinae gen. sp.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                            Sequence 241 from Patent AX056597
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Screening for immunostimulatory dna functional modifyers
Patent: WO 0067023-A 20 09-NOV-2000;
CPG Immunopharmaceuticals GmbH (DE); UNIVERSITY OF IOWA RESEARCH
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Sequence 20 from Patent WO0067023.
AX045790
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Screening for immunostimulatory dna functional modifyers
Patent: WO 0067023-A 17 09-NOV-2000;
CPG Immunopharmaceuticals GmbH (DE); UNIVERSITY OF IOWA RESEARCH
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/mol_type="unassigned DNA"
/mol type="unassigned DNA"
/db xref="taxon:32630"
/noTe="synthetic oligonucleotide"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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; Pred. No. 1.8e
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AX104364/c
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                                                                                                                                                   synthetic construct
synthetic construct
artificial sequences.
                                                                                                                                                                                                                       Sequence 556 from Patent AX104364
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                                                                                 Krieg,A.M., Schetter,C. and Vollmer,J.C. Immunostimulatory nucleic acids Patent: WO 0122972-A 556 05-APR-2001; UNIVERSITY OF IOWA RESEARCH FOUNDATION (
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:39108"
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Pred. No. 1.8e+02;
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Matches 16; Conservative
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                                                                                                                                                                       Nanosphere, Inc. (US)
Location/Qualifiers
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Nanoparticles having oligonucleotides attached thereto and uses
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Sequence 55 from Patent WO0151665.
AX196224
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/mol type="unassigned DNA"
/db xref="taxon:32630"
/noTe="random synthetic sequence"
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/mol_type="unassigned DNA"
/db_xref="taxon;32630"
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AX196239
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AX354974
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20 bp
Sequence 838 from Patent WO0197843.
AX355810
AX355810.1 GI:18620478
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Nanoparticles having oligonucleotides attached thereto
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AX354974
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UNIVERSITY OF IOWA RESEARCH FOUNDATION
Location/Qualifiers
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1.1%; Score 14.8; DB 1; Length
Similarity 88.9%; Pred. No. 1.8e+02;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                     /note="Synthetic oligonucleotide~phosphodiester backbone"
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/db_xref="taxon:32630"
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mol_type="unassigned DNA"
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AX440125
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Sequence 55
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UNIVERSITY OF IOWA RESEARCH FOUNDATION (US)
Location/Qualifiers
Nanoparticles therefor
                           Mirkin, C.A., Leteinger, R.L., Mucic, R.C., Storhoff, J.J., Elghanian, R., Taton, T.A., Park, S.J. and Li, Z.
                                                                                    synthetic construct
synthetic construct
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Methods for enhancing antibody-induced cell lysis
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llarity 88.9%;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
/note="Synthetic oligonucleotide~phosphodiester backbone"
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Pred. No. 1.8e+02;
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Sequence 55 from Patent WO0218643.
AX465311
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                                                                                                                             Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J., Elghanian,R., Taton,T.A., Garimella,V., Li,Z. and Park, Nanoparticles having oligonucleotides attached thereto
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                                                                                 Nanosphere,
                                                                                                Patent: WO 0218643-A 55 07-MAR-2002;
                                                                                                                                                                                           artificial sequences.
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 1.8e+02;
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AX465326
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ACCESSION
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AX524879/c
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Sequence 8 :
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Sequence 70 from Patent WO0218643.
AX465326
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                                                                                                                                                                                           Maruyama, T., Ishiguro, T. and Taya, T. Oligonucleotide and method for detecting Patent: EP 1236806-A 8 04-SEP-2002;
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                                                                                      /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="Oligonucleotide capable of binding
VT2 RNA"
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/db_xref="taxon:32630"
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AX547087/c
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AX547421
LOCUS
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AX547417/c
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Location/Qualifiers
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Similarity 88.9%; Pred. No. 1.8e+02;
16; Conservative 0; Mismatches 2;
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/mol type="unassigned DNA"
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Matches 16; Conserv
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Sequence 55
AX556124
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Sequence 70 from Patent WO0246472.
AX556139
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                                      Mirkin, C.A., Letsinger, R.L., Mucic, R.C., Storhoff, J.J., Elghanian, R., Taton, T.A., Garimella, V., Li, Z. and Park, S.J. Nanoparticles having oligonucleotides attached thereto and
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Location/Qualifiers
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Patent: WO 0246472-A 70 13-JUN-2002; Nanosphere, Inc. (US)
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/db_xref="taxon:32630"
/note="Synthetic Sequence"
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/db_xref="taxon:32630"
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AX664307
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Query Match
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                                                                                                                         chromatography
Patent: WO 0246398-A 6 13-JUN-2002;
The University of Houston System (University of Houston System)
                                                                                                                                                                                            Willson, R.C. and Murphy, J.C. Nucleic acid separation using immobilized
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AX664308
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Location/Qualifiers
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Patent: WO 0246398-A 5 13-JUN-2002;
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                                                                /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
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                                                 /note="Synthetic Oligonucleotide Sequence"
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Pred. No. 1.8e+02;
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AX708893
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                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 334
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TITLE
                                                                     Query Match
Best Local Similarity
Matches 16; Conserv
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Sequence 75
AX708893
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                                                                                                                                                                                           Kirtsen,N.V., Hyldig-Nielsen,J.J. and Williams,B.F.
Methods, kits and compositions pertaining to the suppression of detectable probe binding to randomly distributed repeat sequences in genomic nucleic acid
Patent: WO 03027328-A 14 03-APR-2003;
Boston Probes, Inc. (US); DakoCytomation Denmark A/S (DK)
Location/Qualifiers
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Sequence 14 from Patent WO03027328.
AX741040
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Location/Qualifiers
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ilarity 88.9%;
Conservative
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                                                                                                                       Oligomer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="synthetic construct"
/mol_type="unassigned DNA"
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                                                                                                                               /mol_type="genomic DNA"
/db_xref="taxon:32630"
/note="Description of Combined DNA/RNA Molecule:Synthetic
                                                                                                                                                                      organism="synthetic construct"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note="Oligonucleotide primer"
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Pred. No. 1.8e+02;
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AX741052
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1520 AAAAAAAAAAAGTAAAA 1537
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GENESIS RESEARCH & DEVELOPMENT CO LTD
OS Unidentified
PN UP 2001503969-A/26
PD 27-MAR-2001
PF 28-AUG-1997 JP 1000001157
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Sequence 26 from Patent WO03027328
AX741052
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Prestidge,R.L.
Compounds and methods for treatment and diagnosis of Mycobacterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP 2001503969-A/26. unidentified
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                                                                                                                                                                    PAUL TAN, JUN HIYAMA, ELIZABETH S VISSER, MARGOT A SKINNER, AM SCOTT,
ROSS L PRESTIDGE
A61K39/04, A61K35/74, C07K14/35, C12N15/63
Strandedness: Single;
                                                                                                                                                                                                                                                                                          Unidentified
JP 2001503969-A/26
27-MAR-2001
28-AUG-1997 JP 1998511516
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                                                                                                                                                      opology: Linear
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/note="Description of Combined DNA/RNA Molecule:Synthetic Oligomer Sequence~Synthetic Probe Sequence"
                                                                         /organism='Unidentified'
/organism="unidentified"
/mol_type="genomic DNA"
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Pred. No. 1.8e+02;
0; Mismatches 2;
                                                                                                                                    Location/Qualifiers
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BD107450/c
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BD080522/c
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TITLE
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Best Local Similarity
Matches 16; Conserv
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DN BD107450.1 GI:23202268

US JP 2002034599-A/9:

Synthetic construct

NTSM synthetic construct

Artificial sequences.

1 (bases 1 to 20)

ORS Segawa,M., Takarada,H., Aono,T. and Yoshiga,S.

E Method of detecting single base polymorphism

NAL Patent: JP 2002034599-A 9 05-FEB-2002;

TOYOBO CO LTD

OS Artificial Sequence

PN JP 2002034599-A/9

PD 05-FEB-2002

PF 26-JUL-2000 JP 2000225354

PI MASAYA SEGAWA,HIROSHI TAKARADA,TOSHIYA AONO,SATOKO YOSHIGA PC

C1201/68,C12N15/09,C12N15/09

CC Description of Artificial Sequence:primer

FH Key

FT source 1.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         artificial sequences.

E 1 (bases 1 to 20)

S Pitsch, S., Weiss, P.A. and Jenny, L.
Ribonucleoside-derivative and method for preparing the sam
Patent: JP 2001515087-A 1 18-SEP-2001,
STEFAN PITSCH, PATRICK A WEISS, LUZI JENNY
OS Artificial Sequence
PN JP 2001515087-A/1
PD 18-SEP-2001
PD 18-SEP-2001
PD 18-SEP-2001
PR 18-AUG-1998 JP 2000509723
PR 18-AUG-1998 JP 200509723
PR 18-AUG-1997 CH 1931/97
PI STEFAN PITSCH, PATRICK A WEISS, LUZI JENNY
PO CO7H19/06, CO7F7/18, CO7H19/16, CO7H21/02, CO7H23/00 CC
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D 18-SEP-2001
P 17-AUG-1998 JP 2000509723
PR 18-AUG-1997 CH 1931/97
PI STEFAN PITSCH, PATRICK A WEISS, LUZI JENNY
PC C07H19/06,C07F7/18,C07H19/16,C07H21/02,C07H23/00 CC
Description of Artificial Sequence:synthetic polynucleotic Description of Artificial Sequence:synthetic polynucleotic Description of Artificial Sequence:
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JP 2001515087-A/1.
synthetic construct
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Ribonucleoside-derivative
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ilarity 88.9%;
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/mol_type="genomic RNA"
/db_xref="taxon:32630"
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Pred. No. 1.8e
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AUTHORS
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SOURCE
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AR080294/c
LOCUS
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VERSION
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SOURCE
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BD174543/c
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Sequence 13
AR080294
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synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 20)
Maruyama, T., Ishiguro, T. and Taya, T.
Oligonucleotide for detecting Vero toxin
Patent: JP 2002253257-A 8 10-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PI TAKAHIRO MARUYAMA, TAKAHIKO ISHIGURO, TOSHITAKA TAYA PC C12N15/09, C12Q1/68, G01N33/53, G01N33/566, C12N15/00 CC Oligonucleotide capable of binding specifically to VT2 RNA
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Oligonucleotide for detecting
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                                               1 (bases 1 to 21)
Watson, M.A. and Fleming, T.P.
Mammaglobin, a mammary-specific breast
Patent: US 5968754-A 13 19-OCT-1999;
                                                                                                                          Unknown.
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                                                                                                                                        Unknown.
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Similarity 88.9%;
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JP 2002253257-A/8
10-SEP-2002
02-MAR-2001 JP 2001058143
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                         /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA
/db_xref="taxon:32630"
      /organism≃"unknown"
                                 Location/Qualifiers
                                                                                                                                                                     GI:10007029
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Pred. No. 1
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Pred. No. 1.8e+02
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5968754.
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AR093143/c
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AUTHORS
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FEATURES
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AUTHORS
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AR084524/c
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AR084521
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Best Local Similarity 88.9%;
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16; Conservat
 Sequence 12 from patent US
AR093143
AR093143.1 GI:10019895
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Sequence 13 from patent US
AR084524
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Sequence 10 from patent US
AR084521
                                                                                                                                                                                                                                   1 (bases 1 to 21)
Matson,R.S., Coassin,P.J., Rampal,J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 13 09-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 21)
Matson, R.S., Coassin, P.J., Rampal, J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 10 09-NOV-1999;
                                   AR093143
                                                                                                                                                                                                                                                                                               Unknown
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Similarity 88.9%;
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                                                                                            AAAAAAAAAAAAAAAAA 4
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/mol_type="unassigned DNA"
                                                                                                                                                                                     organism="unknown"
/mol_type="unassigned DNA"
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                                                                                                                                                     Score 14.8; DB 1;
Pred. No. 1.7e+02;
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Pred. No. 1.7e+02;
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Pred. No. 1.7e+02;
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TITLE
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AUTHORS
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AR095412/c
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JOURNAL
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Best Local
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AR095412
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AR103542
                                                                    1 (bases 1 to 21)
Brooks-Wilson,A.R., Buckler,A., Cardon,L.,
Miller,A. and North,M.
Asthma related genes
Patent: US 6087485-A 66 11-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 21)
Watson,M.A. and Fleming,T.P.
Method for detecting the presence of breast cancer increase in mammaglobin mRNA expression
Patent: US 6004756-A 13 21-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 21)
Bergan,R. and Neckers,L.
Inhibition of protein kinase activity
                                                                                                                                           Unknown
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Patent: US 5998596-A 12 07-DEC-1999;
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Similarity 88.9%;
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/mol_type="unassigned DNA"
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                       _type="unassigned
   1.1%;
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Pred. No. 1.7e.
0; Mismatches
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Pred. No. 1.7e+02;
0; Mismatches 2;
   Score 14.8;
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AR153849
LOCUS
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AUTHORS
TITLE
JOURNAL
FEATURES
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BD224108/c
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AR118155/c
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TITLE
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Best Local Similarity
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Mammaglobin, breast cancer BBD224108
BD224108:1 GI:33033878
JP 2002525998-A/10.
Synthetic construct
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Heller, M.J., Tu, E., Evans, G.A. and
Methods for transport in molecular
diagnostics
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AR118155
                                                                                                                                                                                                                                                                                            Patent: US 6238624-A 2 29-MAY-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                           Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2 from patent US AR153849
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Unclassified.
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                                                                                                                                                                                                                                                                organism="unknown"
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pred. No. 1.7e+02;
0; Mismatches 2;
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Pred. No. 1.7e+C
O; Mismatches
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                                                      NA linear PAT 17-JUL-2003 protein specific to mamma.
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1 (Dases 1 to 21)

3 1 (Dases 1 to 21)

5 Watson,M.A. and Fleming,T.P.

5 Watson,M.A. and Fleming,T.P.

6 Mammaglobin, breast cancer secretory protein specific to Patent: JP 2002525098-A 10 13-AUG-2002;

6 WASHINGTON UNIVERSITY

7 OS Artificial Sequence
PN JP 2002525098-A/10
PD 13-AUG-2002
PD 13-AUG-2002
PF 29-SEP-1999 JP 2000572241
PR 29-SEP-1998 US 09/162622
PR 29-SEP-1998 US 09/162622
PR MARK A WATSON,TIMOTHY P FLEMING
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Sequence 2
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136166.1
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Sequence 46 from Patent WO2004036221.
CQ846797 CQ846797.1 GI:50895947
                                                                                                                                                                     16;
                                                                                                                                                                                                                                                                                                                                    O'Toole, M.M. and Liu, W. Compositions and methods for diagnosing and treating autoimmune
                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                             Patent: WO 2004036221-A 46 29-APR-2004; Wyeth (US); O'Toole, Margot Mary (US);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C12N15/09, C12Q1/68, G01N33/53, G01N33/566, G01N33/577//G01N33/574,
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Key Location/Qualifiers
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                                                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="unassigned RNA"
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/db_xref="taxon:32630"
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     GI:2086679
                              from patent US
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Pred. No. 1.7e+02;
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Pred. No. 1.7e+02
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I84433/c
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                                                                          Massively parallel sequencing of sorted Patent: US 5695934-A 23 09-DEC-1997;
                                                                                             1 (bases 1 to 21)
Brenner, S.
                                                                                                                               Unknown.
                                                                                                                                          Unknown.
                                                                                                                                                                         Sequence 23
I84433
                                                                                                                                                                                              I84433
                                                                                                                                                                                                                                                                                                                                                                                 Patent: US 5668267-A 13 16-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                     Polynucleotides encoding mammaglobin,
                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 21)
Watson, M.A. and Fleming, T.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Active programmable electronic devices for molecular biological analysis and diagnostics analysis and Secondary 1987;
Patent: US 5605662-A 2 25-FBB-1997;
Location/Qualifiers
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1 (bases 1 to 21)
Heller, M.J. and Tu, E.
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
                              /organism="unknown"
/mol_type="unassigned DNA"
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Pred. No. 1.
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Pred. No. 1.7
Score 14.8; DB 1;
Pred. No. 1.7e+02;
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AX104720/c
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Watson,M.A. and Fleming,T.P.
Watson,bin, a secreted mammary-specific breast cancer protein
Patent: US 6677428-A 13 13-JAN-2004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                At bases 1 to 21) / Watson, M.A. and Fleming, T.P. Matson, M.A. and Fleming, T.P. Mammaglobin, a secreted mammary-specific breast cancer protein Patent: US 6566072-A 13 20-MAY-2003; Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unknown
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                                                                                                                                                                                                                                                               /mol_type="mRNA"
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                                                                                                                                                                                                                                                                                 /organism="unknown"
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 sequences
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from patent US 6566072.
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88.9%; Pred. No. 1.70
rative 0; Mismatches
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Sequence 840 from Patent
AX355812
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                                                    Inhibition of angiogenesis by nucleic acids Patent: WO 02053141-A 912 11-JUL-2002; Coley Pharmaceutical Group, Inc. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent: WO 0197843-A 840 27-DEC-2001;
UNIVERSITY OF IOWA RESEARCH FOUNDATION
LOCATION/Qualifiers
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Methods for enhancing antibody-induced cell lysis and treating
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Pred. No. 1.7e+02;
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Bookenkamp, D., Dieck, T.H. and Hoppe, H.U. Method for sorting single-stranded nucle Patent: WO 03072818-A 21 04-SEP-2003; Degussa Bioactives GmbH (DE)
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Sequenz:Capture-Oligonukleotid"
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Sequence 23 from Patent WO03072818.
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/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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Sequenz:Capture-Oligonukleotid"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                         /note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
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Pred. No. 1.7e+02;
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Sequence 25 from Patent W003072818.
AX825127
AX825127.1 GI:39750856
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Sequence 26 from Patent WO03072818.
AX825128
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                                                                        Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids Patent: WO 03072818-A 26 04-SEP-2003; Degussa Bioactives GmbH (DE)
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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Pred. No. 1.
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Sequence 27 from Patent W003072818.
AX825129
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mol_type="unassigned DNA"

/db xref="taxon:32630"

/note="Beschreibung der kuenstlichen
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Sequence 29 from Patent WO03072818.
AX825131
AX825131.1 GI:39750860
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Sequence 28 from Patent WO03072818.
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ilarity 88.9%;
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/db_xref="taxon;32630"
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Sequence 30 from Patent WOO3072818.
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AX825132.1 GI:39750861
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/db_xref="texxon:32630"
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/db xref="taxon:35630"
/noTe="Beschreibung der kuenstlichen
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88.9%; Pred. No. 1.70
tive 0; Mismatches
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Sequence 31 from Patent WO03072818.
AX825133
AX825133.1 GI:39750862
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ilarity 88.9%;
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mod_base=OTHER
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                                                Score 14.8; DB 1;
Pred. No. 1.7e+02;
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AX825134/c
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Sequence 32 from Patent WO03072818.
AX825134
AX825134.1 GI:39750863
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Sequence 37 from Patent WO03072818.
AX825139
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Location/Qualifiers
                                                                                        Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic Patent: WO 03072818-A 37 04-SEP-2003; Degussa Bioactives GmbH (DE)
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/organism="synthetic construct"
/mol type="unassigned DNA"
/mol xpef="taxon:32630"
/noTe="Beschreibung der kuenstl
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/mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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/mol_type="unassigned DNA"
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/mod_base=OTHER
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mod_base=OTHER
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RESULT 371
AX825140/c
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Sequence 38 from Patent WO03072818.
AX825140
AX825140.1 GI:39750869
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
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                                                              Nucleic Acid) "
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AX825141/c
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synthetic construct synthetic construct artificial sequences.
                                                                   21 bp
Sequence 40 from Patent W003072818.
XX825142
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Location/Qualifiers
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Sequence 39 from Patent WO03072818.
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/mal_type="unassigned DNA"
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/noTe="Beschreibung der kuenstlichen
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Sequence 41 from Patent WO03072818.
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Sequenz:Capture-Oligonukleotid"
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Sequenz:Capture-Oligonukleotid"
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/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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Sequence 44 from Patent WO03072818
AX825146
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Sequence 43 from Patent WO03072818.
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Sequence 45 from Patent W003072818.
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                /note="LNA-T (Locked Nucleic Acid)"
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/mol_type="taxon:32630"
/noTe="Beschreibung der kuenstl
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Sequence 46 from Patent WO03072818.
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AX825149
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seguenz:Capture-Oligonukleotid"
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Sequence 48 from Patent
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Sequenz:Capture-Oligonukleotid"
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/note="Beschreibung der kuenstlichen
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/mod_base=OTHER
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Pred. No. 1.7e+02;
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                                                                                                   Query Match
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Matches 16; Conserv
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Sequence 54 from Patent WO03072818.
AX825156
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ilarity 88.9%;
Conservative
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/mod_base=OTHER
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/mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                 /note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
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mod_base=OTHER
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Pred. No. 1.7e+02
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Pred. No. 1.7e+02;
0; Mismatches 2
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AX825158/c
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Sequence 55 from Patent W003072818.
AX825157
                                                                                                               Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic Patent: WO 03072818-A 56 04-SEP-2003; Degussa Bioactives GmbH (DE) Location/Qualifiers
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                                                                                                                                                                                                                                          synthetic construct
synthetic construct
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mal_type="unassigned DNA"
/mb.xref="taxon:35630"
/nb.aref="beschreibung der kuenstlichen
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/mod_base=OTHER
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mod_base=OTHER
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/mol_type="unassigned DNA"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noTe="Beschreibung der kuenstl
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mod_base=OTHER
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AX825160/c
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Sequence 58 from Patent WO03072818.
AX825160
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mod_base=OTHER
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/mod_base=OTHER
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 Score 14.8;
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AX825162/c
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DEFINITION
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AX825161/c
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Best Local Similarity 88.9%; Matches 16; Conservative
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Sequence 60 from Patent WO03072818.
AX825162
AX825162.1 GI:39750891
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Sequence 59 from Patent WO03072818
AX825161
                                                                 synthetic construct synthetic construct
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Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids
                                                   artificial sequences
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="LNA-T (Locked Nucleic Acid)"
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Pred. No. 1.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                            synthetic construct synthetic construct artificial sequences.
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Sequence 62 from Patent WO03072818.
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Bioactives GmbH (DE)
/note="LNA-T (Locked Nucleic Acid)"
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12
                                                                                                                                                                                          /organism="synthetic construct"
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/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                             /note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
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/mod_base=OTHER
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AX825165/c
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Sequence 63 from Patent
AX825165
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synthetic construct
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Similarity 88.9%; Pred. No. 1.7e+02;
16; Conservative 0; Mismatches 2
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/mod_base=OTHER
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/mod_base=OTHER
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/mod_base=OTHER
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/mod_base=OTHER
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/mod_base=OTHER
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Seguenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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02-OCT-2001
18-SEP-1998 JP 2000511779
18-SEP-1997 US 08/933149
MARK A WATSON, TIMOTHY P FLEMING
C12N15/09, A61K35/26, A61K39/00, A61K39/00, A61K39/395, A61K39/395, A61P35/00,
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                note="LNA-T (Locked Nucleic Acid)'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                note="LNA-T (Locked Nucleic Acid)"
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Pred. No. 1.7e+02;
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BD087491
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Best Local
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                          1520 AAAAAAAAAAAAGTAAAA 1537
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1 (bases 1 to 21)

Sosnowski,R.G., Butler,W.F., Tu,E., Nerenberg,M.I., Heller,M.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BD087491 PAT 27-AUG-2002 Self-assembling microelectronic integration system capable of designating self address, compartment device, mechanism, method and operation for molecular biological analysis and diagnosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Self-assembling microelectronic integration system capable of designating self address, compartment device, mechanism, method operation for molecular biological analysis and diagnosis Patent: JP 2001525193-A 2 11-DEC-2001;
                                                                                                                                                                                                                                                                                                                           NERENBERG
                                                                                                                                                                                                                                                                                                                                                                                                                            NANOGEN INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BD087491.1 GI:22633101
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                                                                    Similarity
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AAAAAAAAAAAAAAAAAA 18
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JP 2001525193-A/2
11-DEC-2001
                                                    Conservative
                                                                                                                                                                                                                                      MICHAEL J HELLER, CARL F EDMAN
C12Q1/68, C12N15/09, C12N15/00
Description of Artificial Sequence: Synthesized with u
                                                                                                                                                                                                                                                                                                                                      01-DEC-1998 JP 2000524303
05-DEC-1997 US 08/986065
RONALD G SOSNOWSKI,WILLIAM F BUTLER,EUGENE
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Mammaglobin, a secreted mammary specific breast cancer protein
Mammaglobin, Location/Qualifiers
                                                                                                                                                                                                source
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                                                                                                                                                                                                                           secondary structure
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism='Unidentified'
                                                                                                      /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                                                                                                /organism='Artificial
Location/Qualifiers
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                                                    0; Mismatches
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                                                                   Score 14.8; DB 1
Pred. No. 1.7e+02
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Pred. No. 1.7e+02;
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A79657/c
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BD129772
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                                                 Query Match
Best Local Similarity
Matches 15; Conserv
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Best Local Similarity
Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Asthma-associated gene
Asthma-associated gene
Patent: JP 2002500895-A 62 15-JAN-2002;
AXYS PHARMACEUTICALS INC
SUnidentified
PN JP 2002500895-A/62
PN 15-JAN-2002
PD 15-JAN-2002
PD 15-JAN-1998 JP 2000528715
PI ANGELA R BROOKS WILSON, ALAN BUCKLER, LON
CARDON, ALLSOUN H CARRY,
PI MARCARET GALVIN, ANDREW MILLER, MICHAEL NORTH
PC C1201/68, A01K67/027, C07K14/47, C12N15/09, C12N15/00 CC
Strandedness Single;
CC Topology: Linear;
CC Asthma-associated gene
FH Key Location/Qualifiers
FT source / Organism='Unidentified'.
                                                                                                                                                                                                                                                                                                                                                                                                                                          1225 CTCTAGCTCTTAGCTTCCTC 1244
                           1518 ТТАЛАЛАЛАЛАЛАСТАЛА 1536
 19
                                                                                                                                                                             1 (bases 1 to 19)
Emrich, T. and Leying, H. METHOD OF DETECTING TELOMERASE ACTIVITY
PATENT: WO 9720069-A 6 05-JUN-1997;
BOEHRINGER MANNHEIM GMBH (DE); EMRICH TI
                                                                                                                                                                                                                                                                                                                           A79657 19 bp
Sequence 6 from Patent WO9720069.
A79657
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                                                                                                                                                                                                                                                                      unidentified
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Wilson,A.R.B., Buckler,A.,
Miller,A. and North,M.
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JP 2002500895-A/62.
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1..21
/organism='Unidentified'.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                             /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                 GI:6092611
                                                              1.0%;
78.9%;
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                                                      2; Mismatches
                                                                    Score 14.6; DB 1;
Pred. No. 2.2e+02;
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Pred. No. 1.7e+02;
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                                                                                   DB 1;
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AX825107
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AR147331/c
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AX825107
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                                                                                                                                                                                                                                                                                                                                                                                                                                             synthetic construct synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 19)
Emrich, T., Leying, H., Hinzpeter, M. and Karl, G.
Method of detecting telomerase activity
Patent: US 6221584-A 6 24-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6
AR147331
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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/note="Beschreibung der kuenstl:
/note="Beschreibung der kuenstl:
Sequenz:Capture-Oligonukleotid"
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'mod_base=OTHER
                                     /note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
                                                                                                                        /mod
                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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                                                                             note="INA-T (Locked
                                                                                                                                                                                                                                           bound_moiety="Biotin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 organism="unknown"
               note="LNA-T (Locked Nucleic Acid)"
                                                                                                                                                             mod_base=OTHER
                                                                                                                                                                                                   mod_base=OTHER
                                                                                                                                                                                                                                                                                                                          organism="synthetic construct"
                                                                                                                                  note="LNA-T (Locked
                                                                                                                                                                                                                  note="LNA-T (Locked Nucleic Acid)"
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78.9%; Pred. No. 2
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BD085544

BD085544.1 GI:22631154

BD085544.1 GI:22631154
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Sequence 15
AX825117
AX825117.1
Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 22)
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llarity 81.0%;
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/mod_base=OTHER
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Seguenz:Capture-Oligonukleotid"
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Sequence 5561 from Patent EP1281758.
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AX692829.1 GI:29415792
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                      Unclassified
                                                               Unknown.
                                                                                                      AR089217.1
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JP 2001333800-A/1
04-DEC-2001
30-MAY-2000 JP 2000160324
KAORI SHIWADA
C12Q1/68,C12N15/09,G01N33/50,C12N15/00
Method of comparison and detection of RNA amount and DNA
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                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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/mol_type="genomic RNA"
/db_xref="taxon:9606"
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Pred. No. 1.9e+02
0; Mismatches
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Pred. No. 1.6e+02;
0; Mismatches 4,
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1 (bases 1 to 17)
Pavco, P. A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J. A. Roberts, E. and the conditions concerning molecule participating in vasculogenic response patent: JP 2002509721-A 4537 02-APR-2002; PARRACEUTICALS INC
OS HOMO Sapiens (human)
PN JP 2002509721-A/4537
PD 02-APR-2002
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BD201512
Method and reagent
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DNA constructs and methods for screening for increased expression of human apo AI gene
Patent: US 5994061-A 13 30-NOV-1999;
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BD201511.1 GI:33011281
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27-MAR-1998 US 60/079678
PAMELA A PAVCO, ELISABETH ROBERTS, THALE JARVIS, CLAIRE COESHOTT,
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Key Location/Qualifiers
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                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="genomic RNA"
/db_xref="taxon:9606"
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Location/Qualifiers
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/mol_type="unassigned DNA"
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Pred. No. 2.7e+02;
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Pred. No. 2.7e+02;
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BD258338
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PN JP 200254795-A/6131
DP 10-DEC-2002
PD 11-APR-2000 JP 2000611654
PR 12-APR-1999 US 60/1129390
PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC C12P21/02,
                                                                                                                                                                                             1 (bases 1 to 17)
Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.
Regulation of repressor genes using nucleic acid molecules
Patent: JP 2002541795-A 6131 10-DEC-2002;
RIBOZYME PHARMACEUTICALS INC
                                                                                                                                                                                                                                                                                                                                        BD258338.1 GI:33068108
JP 2002541795-A/6131.
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1 (Dases 1 to 17)

Pavco, P.A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J.A. Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response Patent: JP 2002509721-A 4538 02-APR-2002;
     C12P21/02, C12P21/02//A61K31/711, (C12N5/10, C12R1:91), (C12P21/02, C12R1:91),
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BD201512
BD201512.1 GI:33011282
JP 2002509721-A/4538.
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27-MAR-1998 US 60/079678
PAMELA A PAYCO ELISABETH ROBERTS, THALE JARVIS, CLAIRE COESHOTT,
JAMES A MCSWIGGEN
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Key Location/Qualifiers
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JP 2002509721-A/4538
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/mol_type="genomic RNA"
/db_xref="taxon:9606"
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Pred. No. 2.7e+02;
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acid molecules:
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DEFINITION
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AUTHORS
TITLE
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BD258339
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Best Local Similarity 93.8
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1 (bases 1 to 17)

Blatt,L., Zwick,M., Pavco,P. and Mcswiggen,J.

"E Regulation of repressor genes using nucleic acid

NAL Patent: JP 200541795-A 6132 10-DEC-2002;

RIBOZYME PHARMACEUTICALS INC

OS Eukaryote
PN JP 2002541795-A/6132

PD 10-DEC-2007
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PD 10-DEC-2002

PD 11-APR-2000 JP 2000611654

PF 11-APR-2000 JP 2000611654

PR 12-APR-1999 US 60/129390

PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN

C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC
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 CQ616719
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A61K37/02,
A61K37/02,
(C12N5/00,C12R1:91)
(C12N5/00,C12R1:91)
Regulation of repressor genes using
Location/Qualifiers
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Regulation of repre
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                                                                                                                                                                                                           /organism='Eukaryote'
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                                                                                                                                                            /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Pred. No. 2.7e+02;
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Sequence 1460 from Patent
CQ616720
CQ616720.1 GI:41666938
                                 Homo sapiens (human)
                                                                       CQ617530
Sequence 2270 from Patent
CQ617530
CQ617530.1 GI:41667748
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           Eukaryota; Metazoa;
Mammalia; Eutheria;
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Mammalia; Eutheria;
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Sequence 1459 from Patent W00192524.
CQ616719
CQ616719.1 GI:41666937
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Myosin-like gene expressed in human heart Patent: WO 0192524-A 1460 06-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. and
Shannon,M.E.
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Similarity 93.8%; Pred. No. 2.7e+02;
15; Conservative 0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Primates;
Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 2.7e+02;
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Sequence 2545 from patent
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Sequence 2271 from Patent
CQ617531
CQ617531.1 GI:41667749
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                                                              1 (bases 1 to 17)
Pavco, P., McSwiggen J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 2545 12-FEB-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. and Shannon,M.E.
Myosin-like gene expressed in human heart and muscle
Patent: WO 0192524-A 2271 06-DEC-2001;
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Homo sapiens
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Mammalia; Eutheria; Primates;
                                                                                                                                              Unclassified.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
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                                        organism="unknown"
                         mol_type="unassigned DNA"
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AR397930/c
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AR285940
Sequence 311 from patent
AR397930
AR397930.1 GI:40135323
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Beigelman, L., Burgin, A., Beaudry, A., Karpeisky, J.
Matulic-Adamic, J., Sweedler, D. and Zinnen, S.
Synthetic ribonucleic acids with RNAse activity
Patent: US 6528640-A 312 04-MAR-2003;
                                                                                                                                                                                                                                                    Unclassified.

1 (bases 1 to 17)

Pavco,P., McSwiggen,J.A., Stinchcomb,D.T. and Escobedo,J.

Pavco,P., McSwiggen for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6566127-A 1069 20-MAY-2003;
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/mol_type="unassigned
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93.8%; Pred. No. 2.7e+02;
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AR457783/c
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Sequence 1459 from patent
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                                       1 (bases 1 to 17)
Gu,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Shannon,M.E.
Polynucleotide encoding a human myosin-like polypredominantly in heart and muscle patent: US 6865188-A 1460 03-FEB-2004;
Location/Qualifiers
                                                                                                                                    Unknown.
Unclassified.
                                                                                                                                                                                          AR457783
Sequence 1460 from patent
AR457783
AR457783.1 GI:42692840
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Gu,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. Shannon,M.E.
Polynucleotide encoding a human myosin-like polypeptide e predominantly in heart and muscle predominantly in heart and muscle Patent: US 6686188-A 1459 03-FEB-2004;
Location/Qualifiers
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Unclassified.
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/organism="unknown"
/mol_type="genomic"
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93.8%; Pred. No. 2.7e
tive 0; Mismatches
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                                                                                                            Chen, W.
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Best Local Similarity 93.8%;
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AX674744 17 bp DNA Sequence 3189 from Patent WO03004526.
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AR458593
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Sequence 2271 from patent
AR458594
                                                                                                                                                                                                                 Polynucleotide encoding a human myosin-like polypeptide expressed predominantly in heart and muscle Patent: US 6686188-A 2271 03-FEB-2004;
                                                                                                                                                                                                                                                              1 (bases 1 to 17)
GU.Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R.,
                                                                                                                                                                                                                                                                                                      Unknown.
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GRY, Ji, Y., Penn, S.G., Hanzel, D.K., Rank, D.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unknown.
                                                                                                                     h 1.0%; Score 14.4; DB 1;
Similarity 93.8%; Pred. No. 2.7e+02;
15; Conservative 0; Mismatches 1
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                                                                                                                                                                  /mol_type="genomic DNA"
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                                                                                                                                                                               organism="unknown"
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Pred. No. 2.7e+02;
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AX730083/c
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                                                                                   AX730083
Sequence 1717 from Patent
AX730083
                            Homo sapiens (human)
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Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                     Telerman, A., Amson, R. and Tuijnder, M.
Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as medicines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 1189 from Patent
AX729555
AX729555.1 GI:30508898
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and their use as
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AX674744.1 GI:29333092
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                      AX730083.1 GI:30509426
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Pred. No. 2.7e+02;
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                                    1 (bases 1 to 18)
Brysch, W. and Schlingensiepen, K.
BRYSCH, W. and Schlingensiepen, K.
ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD
PATENTE: WO 9833904-A 1526 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
Location/Qualifiers
                                                                                                                                                                                                                                                                            Sequence 1526 from Patent A89378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in tumoral suppression, tumoral reversion apoptosis and/or viral resistance phenomena and their use as
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Sequence 5017 from Patent
AX761696
AX761696.1 GI:32256312
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Location/Qualifiers
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                                                                                                                                                                    unclassified
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/mol_type="unassigned DNJ
/db_xref="taxon:9606"
/organism="unidentified"
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/db_xref="taxon:9606"
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AX662307/c
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AR106885
                                                                                                                                                                       synthesis
Patent: WO 02059293-A 46 01-AUG-2002;
Patent: Anthony C. (US) ; Blacklow,
                                                                                                                                                                                                                                                    synthetic construct
synthetic construct
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Sequence 46 from Patent
AX662307
AX662307.1 GI:29163190
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Cowsert, L.M., Bennett, C.Frank. and O'Malley, B.W.
Antisense modulation of SRA expression
Patent: US 6107092-A 46 22-AUG-2000;
Location/Qualitiers
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                                                                                                                                                                                                                                         artificial sequences.
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larity 93.8%;
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/mal_type="unassigned DNA"
/mal_type="taxon:32630"
/mbee="from Synthetic DNA"
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/db_xref="taxon:32644"
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Pred. No. 2.5e+02;
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 WO 0142475-A/27.
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          R1420FP1
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S Schlingensiepen, K.H. and Brysch, W.
An antisense oligonucleotide preparation method
L Patent, JP 2001511000-A 1526 07-AUG-2001;
BICGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
OS UDAROWN
P 2001511000-A/1526
PD 07-AUG-2001
PF 30-JAN-1998 JP 1998532533
PF 31-JAN-1997 EP 97101531.8
PF 31-JAN-1997 EP 97101531.8
PI KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
PC C12N15/11, C07H21/04, A61K31/70
CC An antisense oligonucleotide preparation method FH
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Patent: WO 0142475-A 27 14-JUN-2001;
JAPAN AS REPRESENTED BY DIFFERENCE GENERAL OF MINISTRY OF AGRICULTURE
FORESTRY AND FISHERIES NATIONAL INSTITUTE OF AGROBIOLOGICAL
RESOURCES, KENICHI NOGUCHI YUKO OHASHI,ICHIRO MITSUHARA,TAKUJI
SASAKI,YOSHIAKI NAGAMURA, HIROYUKI ITO, TAKAYOSHI IWAI,SUSUMU
                                                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 18)
Ohashi, Y., Mitsuhara, I., Sasaki, T., Nagamura, Y., It, H., Iwai, T. and Hiraga, S.
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                                            HIROYUKI ITO,
TAKAYOSHI IWAI,SUSUMU HIRAGA
C12N15/53,C12N9/08,C12Q1/68
                                                                                                                  14-JUN-2001
08-DEC-2000 WO 2000JP008728
10-DEC-1999 JP 99P 352472
YUKO OHASHI,ICHIRO MITSUHARA,TAKUJI SASAKI,YOSHIAKI NAGAMURA,
                                                                                                                                                                                                                    Oryza sativa (rice)
WO 0142475-A/27
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                         artificial sequences.

1 (bases 1 to 20)

Besterman, J.M., MacLeod, A.R. and Siders, W.M.
Modulation of gene expression by combination the Patent: JP 2002528391-A 47 03-SEP-2002;
METHYLGENE INC
OS Artificial Sequence
PN JP 2002528391-A/47
PD 03-SEP-2002
PF 19-OCT-1999 JP 2000576885
PF 19-OCT-1999 US 60/104804
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GENSET (FR)
Other publication AU 6432094 941024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 (bases 1 to 20)
Vasseur,M., Blumenfeld,M., Meguenni,S. and Poddevin,B.
STAPLE AND SEMI-STAPLE OLIGONUCLECTIDES, METHOD OF PREPARATION AND APPLICATIONS
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Sequence 5 from Patent WO9423026.
A40129
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PF 19-OCT-1999 JP 2000576885
PR 19-OCT-1998 US 60/104804
PI JEFFREY M BESTERMAN, ALAN ROBERT MACLEOD, WILLIAM M SIDERS PC A61K48/00, A61K31/165, A61K31/19, A61K31/513, A61K31/517, A61K31/ PC
                                                                                                                                                                                                                                BD244919.1 GI:33054689 JP 2002528391-A/47.
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Modulation of gene expression
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="genomic DNA"
/db_xref="taxon:4530"
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AR298254
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                                                                   l (bases 1 to 20)
Griffals,R., Hoiseth,S.K., Zagursky,R.J.,
Sankaran,B. and Fletcher,L.D.
Chlamydia pneumoniae polynucleotides and u
Patent: US 6559294-A 6476 06-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 20)
Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing
disequilibrium map of the human genome
Patent: US 6537751-A 9989 25-MAR-2003;
                                                                                                                                                                                          Sequence 6476 from AR315939
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                    /mol_type="genomic DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                   /organism="unknown"
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Sequence 114 from patent
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1 (kmets,R., Anderson,K.L., Dean,M., Leppert,M., Lewis,R.A.,
Li,Y., Luppeki,J.R., Nathans,J., Rattner,A., Shroyer,N.F., Singh,N.,
Smallwood,P. and Sun,H.
Smallwood,P. and sun, acid sequences for ATP-binding cassette
transporter and methods of screening for agents that modify
ATP-binding cassette transporter
patent: US 6713300-A 114 30-MAR-2004;
Location/Qualifiers
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                                                                                                                                                                                                    Boekenkamp,D., Hoppe,H.U. and Burgstaller,P. Detection system for separating constituents production and use of the same Patent: WO 0071747-A 35 30-NOV-2000; Aventis Research & Technologies GmbH & Co. KC
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Sequenz:Erkennungssystem"
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                                                                                                                                   organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="genomic I
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Sequence 6 from Patent WO0071703.
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Sequence 2046 from Patent
AX184293
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Sequence 15 from Patent
AX053091
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                                                                                                                                                                                                                                                                                                                                                                                                                                     synthetic construct artificial sequences.
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Inhibition of histone deacetylase
Patent: WO 0071703-A 6 30-NOV-2000;
Methylgene, Inc. (CA)
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                                                                                                                                                                                                                 1.0%;
Similarity 93.8%;
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sapiens
sapiens
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                                                                                                                                                                                                                                                       /note="Description of Combined DNA/RNA Molecule: Positions 1-4 and 17-20 are 2'-methoxyribose substituted nucleotides; positions 5-16 are deoxyribonucleotides"
                                                                                                                                                                                                                                                                                               /organism="synthetic construct"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noTe="synthetic oligonucleotide"
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Pred. No. 2.3e+02;
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WO0142511.
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Sequence 51
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Patent: WO 0142511-A 2046 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
Biotherapeutics Corporation (CA)
                                                                    Besterman, J.M., Macleod, A.R. and Siders, W.M. Modulation of gene expression by combination Patent: EP 1243289-A 51 25-SEP-2002;
                                                                                                                                            synthetic construct synthetic construct
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                       Methylgene, Inc.
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larity 88.2%;
Conservative
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 /organism="synthetic construct"
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/mol_type="unassigned
                                        Location/Qualifiers
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Pred. No. 2.3e+02;
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Pred. No. 2.3e+02;
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                                                                                     therapy
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A90381/c
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A88414/c
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A88414
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Brysch,W.D. and Schlingensiepen,K.D.
An antisense oligonucleotide preparation method Patent: EP 0856579-A 562 05-AUG-1998;
                                                                                                                                                                                                                                                                Sequence 562 from Pat
A90381
A90381.1 GI:6738895
                                                                                                                                                                                                                                                                                                                                                                                                                16;
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Brysch, W. and Schlingensiepen, K.
ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD
Patent: WO 9833904-A 562 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
Location/Qualifiers
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                                                                                                                                          BIOGNOSTIK GES (DE)
Location/Qualifiers
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llarity 84.2%;
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562 from Patent WO9833904.
                                                                                          /organism="unidentified"
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                                                         Score 14.2; DB 1;
Pred. No. 2.7e+02;
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Pred. No. 2.7e+02;
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Pred. No. 2.3e+02;
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PF 2002360271-A/587
PD 17-DEC-2002
PF 28-NOV-2001 JP 2001362595
PF 28-NOV-2001 TW 90110785
PR 04-MAY-2001 TW 90110785
PI CHANG-YEE LING, RUEY-WEN LIN, ZHOU-MENG YOO, XIN-HSUAN HUANG, BE HARING LEE, YI-JU LIN, CI-CHUNG HUANG, HAN-CHANG HSU, CHANG HSI, WEN SHI, PI CHIN-XIN YEH, YI-FENG CAO, CHIH-LONG PAN PI C12N15/09, C12N15/09, C12N1/34, C12Q1/04, C12Q1/42, C12Q1/68 PC (C12N15/09, C12N13/574, G01N33/58, G01N37/00// (C12M1/34, C12R1:9)
PC G01N33/53, G01N33/574, G01N33/58, G01N37/00// (C12M1/34, C12R1:9)
PC C12Q1/70, C01XR1:93), C12N15/00, C12N15/00
CC Oligonuclectide M065108 for identifying HPV CP8061. FH Key
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Sequence
AR294423
                                                                                                                      1 (bases 1 to 19)
Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing disequilibrium map of the human genome Patent: US 6537751-A 6158 25-MAR-2003; Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  artificial sequences.

1 (bases 1 to 19)

Ling, C., Lin, R., Yoo, Z., Huang, X., Lee, B., Lee, S., Lin, Y., Huang, C., Hsu, H., Shi, C., Yeh, C., Cao, Y. and Pan, C.

Method and detector for identifying subtypes of human papiloma Patent: JP 2002360271-A 587 17-DEC-2002;

KING CAR FOOD INDUSTRIAL CO LTD

OS Artificial Sequence
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Method and detector for identifying subtypes of human papiloma
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(C12Q1/70,C12R1:93),C12N15/00,C12N15/00
Oligonucleotide M806108 for identifying HPV CP8061. FH Key
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28-NOV-2001 JP 2001362595
04-MAY-2001 TW 90110785
CHING-YEE LING, RUEY-WEN LIN, ZHOU-MENG YOO, XIN-HSUAN HUANG, BOW-HAENG LEE,
   Conservative
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Score 14.2; DB 1;
Pred. No. 2.7e+02;
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Ribozyme therapy for the treatment of proliferative
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db xref="taxon:32630"
/note="position 65"
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/mbl_tyse="taxon:9606"
/mbe="PCNA HH ribozyme binding
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Pred. No. 2.7e+02;
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BD065927/c
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JP 2001511000-A/562.
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Sequence 558 from Patent
AX742755
AX742755.1 GI:30576744
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Schlingensiepen, K.H.
                                    unidentified
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King Car Food Industrial Co., Ltd. (T
Location/Qualifiers
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Method and detector for identifying subtypes of human papilloma
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and Brysch, W.
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Pred. No. 2.7e+02;
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Sequence 57 from Patent WO9813490.
A70736
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Other publication FR 2703053 940930
Location/Qualifiers
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Vasseur,M., Blumenfeld,M., Meguenni,S. and Poddevin,B.
STAPLE AND SEMI-STAPLE OLIGONUCLECTIDES, METHOD OF PREPARATION AND APPLICATIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            An antisense oligonucleotide preparation method
Patent: JP 2001511000-A 562 07-AUG-2001;
BIOGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
unidentified
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llarity 84.2%;
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KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
C12N15/11, C07H21/04, A61K31/70
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07-AUG-2001
30-JAN-1998 JP 1998532533
31-JAN-1997 EP 97101531
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Pred. No. 2.7e+02;
0; Mismatches 3;
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Pred. No. 2.5e+02;
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Best Local Similarity 84.2%;
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Sequence 57 from Patent EP0834561.
A79220
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 l Similarity
16; Conservat
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/mol_type="unassigned DNA"
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                                      /organism="unknown"
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 Score 14.2; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 3;
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Pred. No. 2.5e+02;
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AR086311/c
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AR129739
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ARI29739
ARI29739.1 GI:14117636
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Dean, N.M., McKay, R., Miraglia, L. and I
Antisense oligonucleotide compositions
inibition of c-Jun and c-Pos
Patent: US 598558-A 132 16-NOV-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                1 (bases 1 to 20)
MCKAY,R., Butler,M.M., Wyatt,J. and Cowsert,L.M.
Antisense modulation of pepck-cytosolic expression
Patent: US 6187545-A 143 13-FEB-2001;
Location/Qualifiers
                       Unknown
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AR142677
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/mol_type="unassigned
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Pred. No. 2.5e+02;
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Sequence
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1 (bases 1 to 20)
Shannon, K.W., Wolber, P.K., Delenstarr, G.C., Kincaid, R.H.
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Sequence 340 from patent
AR158718
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                                                                                                                                          Unclassified.
1 (Dases 1 to 20)
Shannon, K.W., Wolber, P.K., Delenstarr, G.C., Kincaid, R.H.
Method for evaluating oligonucleotide probe Patent: US 6251588-A 340 26-JUN-2001;
                                                                                                                                                                                                                                                     AR158718.1
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                                                          Similarity
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Sequence 12:
AR176870
AR176870.1
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Sequence 6
AR169510
AR169510.1
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AR163731
                                                                                                                                                                                                                                16;
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Bartel, P.L. and Tavtigian, S.V.
MMSC2--an MWAC1 interacting protein
Patent: US 6291173-A 6 18-SEP-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 20)
Bennett, C. Frank. and Cowsert, L.M.
Antisense inhibition of cytchesin-2
Patent: US 6271029-A 18 07-AUG-2001;
Location/Qualifiers
Unclassified.
1 (Dases 1 to 20)
Denn,N.M., McKay,R., Miraglia,L. and Baker, Antisense oligonucleotide compositions and modulation of activating protein 1
Patent: US 6312900-A 125 06-NOV-2001;
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llarity 84.2%;
Conservative
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                                                                                                                                                                                                                                                                      /mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        organism="unknown"
                                                                                                                                                                                                                                                                                 organism="unknown"
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Pred. No. 2.5e
0; Mismatches
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2.5e+02;
les 3;
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linear

PAT 18-JUN-2001

Indels

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RESULT 459
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AR176877/c
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Best Local Similarity
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                     Patent: JP 1993244995-A 7 2-
KYOWA HAKKO KOGYO CO LTD
OS Artificial gene
OC Artificial sequence; Ge
OS Zea maize
PN JP 199324995-A/7
PD 24-SEP-1993
PF KOMATSU YUKI, KIKUCHI
PC C1201/68 C12N15/11;
CC strandedness: Single;
CC topology: Linear;
CC hypothetical: No;
CC anti-sense: No.
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Dean,N.M., McKay,R., Miraglia,L. and Baker,B.
Antisense oligonucleotide compositions and methods for the modulation of activating protein 1
Patent: US 6312900-A 132 06-NOV-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 132 from patent AR176877
                                                                                                                                                                                                                                             synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 20)
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16; Conservative
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                                                                           Zea maize
JP 1993244995-A/7
24-SEP-1993
24-SEP-1991 JP 1991243122
KOMATSU YUKI, KIKUCHI YASUHIRO
C12Q1/68,C12N15/11;
                                                                                                                                                                                                                                                                                                                                                          primer for detecting
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    and Kikuchi, Y.

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/mol_type="unassigned DNA"
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            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                    GI:2173685
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84.2%; Pred. No. 2.5e
ative 0; Mismatches
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Pred. No. 2.5e
0; Mismatches
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AUTHORS
TITLE
JOURNAL
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E28096/c
LOCUS
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ORGANISM
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DEFINITION
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E44159/c
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Best Local
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Best Local Similarity 84.2%;
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CC Topology: Linear;
FH Key
FT source
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                                                                                                                                                                          E44159 20 bp DNA linear PAT 27-AUG-2002 Methods of identification and specific detection of slow-growing mycobacteria by using characteristic base sequence occurring in DNA
                                                                                                                                                                                                                                                                                                                                  16;
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HidekI,K. and Senshu,U.
Method for analyzing DNA fragment
Patent: JP 1999196874-A 7 27-JUL-1999;
HITACHI LTD
gyrase gene
Patent: JP (
                     mycobacteria by using characteristic base sequence occurring in DNA
                                     artificial sequences.

1 (bases 1 to 20)

Kasai,H., Ezaki,T. and Harayama,S.
Methods of identification and specific detection of slow-growing
                                                                                           synthetic construct
synthetic construct
                                                                                                                                                   gyrase gene.
E44159
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16; Conservative
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C12N15/09,C12Q1/68,G01N27/447,C12N15/00,G01N27/26 CC
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JP 1999196874-A/7
27-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Topology: Linear;
Key
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for analyzing DNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism='Unidentified'.
                                                                                                                                                                                                                                                                                                                                                                                 /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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2001128679-A 2 15-MAY-2001;
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84.2%; Pred. No. 2.5e+02;
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Pred. No. 2.5e+02;
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Indels

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Gaps

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Gaps

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PAT 07-OCT-1997

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RESULT 463
117092
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AUTHORS
TITLE
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VERSION
KEYWORDS
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E59328
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Best Local Similarity
Matches 16; Conserv
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Best Local
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Patent: JP 2000342265-A 9 12-DEC-2000;
POAGOSEI CHEM IND CO LTD
OS Artificial Sequence
PN JP 2000342265-A/9
PD 12-DEC-2000
PF 02-UN-1999 JP 1999154974
PR 02-UN-1999 JP 1999154974
PR KUNIHIKO HIROSE, TADAO YOSHIDA
PC C12N15/09, B01D15/08, C12N15/00
CC Key Location/Qualifiers
FT source 1, 20
FT source 1, 20
FT formal forma
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synthetic construct
artificial sequences
1 (bases 1 to 20)
1 Hirose, K. and Yoshida, T.
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OS Artificial Sequence
PN 72001128679-A/2
PD 15-MAY-2001
PF 02-NOV-1999 JP 1999312525
PI HIROAKI KASAI, TAKAYUKI EZAKI, SHIGEAKI HARAYAMA PC C12N15/09, C12Q1/04, C12Q1/68, C12N15/00
CC
FH Key Location/Qualifiers.
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Sequence
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JP 2000342265-A/9.
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Raben, N., Nichols, R., Plotz, P. and Leff, R.
Raben, N., Nichols, R., Plotz, P. and Leff, R.
Assay using recombinant histidyl-tRNA synthetase
Patent: US 5484703-A 7 16-JAN-1996;
Location/Qualifiers
                                                             l Similarity
16; Conservat
                                                                                                                                                                                                I (bases 1 to 20)
Zonana,J., Ferguson,B.M., Headon,D. and Overbeek,P.
Hypohidrotic ectodermal dyplasia genes and proteins
Patent: US 6355782-A 51 12-MAR-2002;
Location/Qualifiers
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Raben,N., Nichols,R., Plotz,P. and Leff,R.
Raben,N., Nichols,R., Plotz,P. and Leff,R.
Assay using recombinant histidyl-tRNA synthetase
Patent: US 5663066-A 7 02-SEP-1997;
Location/Qualifiers
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Unclassified.
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ilarity 84.2%;
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/mol_type="unassigned DNA"
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/mol_type="unassigned
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/mol_type="unassigned
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Unclassified.
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1 (bases 1 to 20)
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3 Bennett, C. F. and Freier, S. M.
Antisense modulation of glioma-associated on/
AL Patent: US 6440739-A 83 27-AUG-2002;
Location/Qualifiers
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Sequence 83
AR224778
AR224778.1
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Sequence 4133 from patent
AR313596
AR313596.1 GI:31707022
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AR307902
AR307902.1 GI:31698658
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Unclassified.

1 (bases 1 to 20)

Griffais,R., Hoiseth,S.K., Zagursky,R.J., Metcalf,B.J., Sankaran,B. and Fletcher,L.D.

Chlamydia pneumoniae polynucleotides and uses thereof
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ilarity 84.2%;
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/mol_type="genomic DNA"
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Pred. No. 2.5e+02;
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Location/Qualifiers
N
                                           16;
                                                                                                                                  Unclassified.
1 (bases 1 to 20)
Buchardt,O., Egholm,M., Nielsen,P.E. and Peptide nucleic acids
Patent: US 6395474-A 4 28-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                   16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unclassified.
1 (bases 1 to 20)
Griffais, R., Hoiseth, S.K., Zagursky, R.J.,
Sankaran, B. and Fletcher, L.D.
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Sequence 5443 from patent
AR314906
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/mol_type="genomic |
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                                          Score 14.2; D
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RESULT 471

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AR428075
 1 (bases 1 to 20)
Buchardt, O., Buchardt, D., Egholm, M., Nielsen, P.E.
Synthetic procedures for peptide nucleic acids
Patent: US 6713602-A 4 30-MAR-2004;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 20)
Buchardt,O., Egholm,M., Nielsen,P.E.
Peptide nucleic acid synthons
Patent: US 6710163-A 4 23-MAR-2004;
Location/Qualifiers
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Spear, P.G., Warner, M.S., Geraghty, R.J., Martinez, W.M.,
Montgomery, R.I., Cohen, G.H., Eisenberg, R.J., Whitbeck, C.J.
Krummenacher, C.
Cellular proteins which mediate herpesvirus entry
Cellular proteins which mediate herpesvirus entry
Patent: Us 6641818-A 5 04-NOV-2003;
Location/Qualifiers
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Sequence 4 from patent US
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/mol_type="genomic DNA"
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Sequence 15 from Patent WO0105435.
AX078001 GI:13157746
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/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref≈"taxon:9606"
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/mol_type="genomic
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Antisense therapy for hormone-regulated tumors Patent: WO 0105435-A 15 25-JAN-2001; THE UNIVERSITY OF BRITISH COLUMBIA (CA); Miyal Location/Qualifiers
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                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                                                                                                                                                                                                                                                 synthetic construct synthetic construct
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Identification method and specific detection
mycobacteria utilizing dna gyrase gene
Patent: EP 1098003-A 3 09-MAY-2001;
MARINE BIOTECHNOLOGY INSTITUTE CO., LTD. (JP)
Location/Qualifiers
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synthetic construct
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Sequence 3 from Patent EP1098003.
AX137428 GI:14273633
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Pred. No. 2.5e+02;
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Pred. No. 2.5e+02;
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AX462672/c
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AX452909/c
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                                                                                   SOURCE
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                            Aegilops tauschii
Aegilops tauschii
Bukaryota, Viridiplantae, Streptophyta; Em
Spermatophyta; Magnoliophyta; Liliopsida;
Pooideae; Triticeae; Aegilops.
Bernard,M., Sourdille,P. and Guyomarch,H. Microsatellite markers from Triticum tauschii Patent: EP 1217079-A 416 26-JUN-2002;
                                                                                                                                                                                                                                Sequence 416 from Patent AX462672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jackson,D., Casari,G. and Suckow,J. Mammalian nuclear receptor cofactors cf7 Patent: WO 0242322-A 10 30-WAY-2002; LION Bioscience AG (DE)
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Similarity 84.2%;
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/mol_type="unassigned DNA"
/db_xref="taxon:10095"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Pred. No. 2.5e+02;
0; Mismatches 3
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Pred. No. 2.5e+02;
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a; Poales; Poaceae;
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AX750458/c
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AX462737/c
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Best Local Similarity
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16; Conserv
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Microsatellite markers from Triticum tauschii
Patent: EP 1217079-A 481 26-UUN-2002;
INSTITUT NATIONAL DE LA RECHERCHE AGRONOMIQUE (INRA)
LOCALION/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Aegilops tauschii
Aegilops tauschii
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Pooideae; Triticeae; Aegilops.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 481 from Patent AX462737
                                                                                                                                                          Isogai,T., Sugiyama,T., Otsuki,T., Wakamatsu,A., Sato,H., Ishii,S.,
Yamamoto,J.I., Isono,Y., Hio,Y., Otsuka,K., Nagai,K., Irie,R.,
Tamechika,I., Seki,N., Yoshikawa,T., Otsuka,M., Nagahari,K. and
                                                                                                                                                                                                                                     synthetic construct
synthetic construct
                                                                                                                                                                                                                                                                              AX750458
Sequence 3983 from Patent
AX750458
AX750458.1 GI:32132876
                                                                                                                  Patent: EP 1308459-A 3983 07-MAY-2003;
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                                                                                                 Helix Research Institute (JP) ; Research
                                                                                                                                Full-length cDNA sequences
                                                                                                                                             Masuho, Y.
                                                                                                                                                                                                                       artificial sequences.
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            /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Aegilops tauschii"
/mol_type="unassigned DNA"
/db_xref="taxon:37682"
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/db_xref="taxon:37682"
                                                                       Location/Qualifiers
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note="an artificially synthesized primer sequence"
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Pred. No. 2.5e+02;
0; Mismatches 3
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Pred. No. 2.5e+02;
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AX785487/c
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Susceptibility locus for schizophrenia
Susceptibility locus for schizophrenia
Susceptibility locus for schizophrenia
Gurling, Hugh Malcolm Douglas (GB)
Gurling, Hugh Malcolm Townsiderars
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                                                                                                                                                                                                                                                                                           Oreochromis niloticus (Nile tilapia)
Oreochromis niloticus
Oreochromis niloticus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Euteleostei; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes;
Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes;
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                                                                                                                                                                                             Genomar ASA (NO)
                                                                                                                                                                                                                                     Lie,Y., Slettan,A., Hoeyum,M. and Lingaas,F. Verification of food origin based on nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     artificial sequences.
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                                                                                                                                                                                                    recognition
Patent: WO 03060160-A 1139 24-JUL-2003;
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                                                                         Similarity
                                                                                                                                                                                                                                                                               abroidei; Cichlidae; Oreochromis.
                                TGTTGCTGCTGCCTTTGTG 747
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                                                           Conservative
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                                                                                                                  organism="Oreochromis niloticus"
|mol_type="unassigned DNA"
|db_xref="taxon:8128"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mol_type="unassigned
/db_xref="taxon:32630"
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                                                                                                                                                                           ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note="Primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="synthetic construct"/mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                      GI:38522112
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                                                                       1.0%;
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Pred. No. 2.5e+02;
0; Mismatches 3;
                                                           Score 14.2; D
Pred. No. 2.5e
0; Mismatches
                                                           0
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Pred. No. 2
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BD056369
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Bource
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                                                                                                                                                                                                                                                                                                                          486
                synthetic construct

artificial sequences.

CE 1 (bases 1 to 20)

RS Takagi.M., Shinji.H. and Ota,K.

RS Takagi.M., Shinji.H. and Ota,K.

Peptide having a function regulating transcription of gene

Patient: JP 2001269176-A 3 02-OCT-2001;

AGENCY OF IND SCIENCE & TECHNOL

OS Artificial Sequence

PN JP 2001269176-A/3

PD 02-OCT-2001

PF 27-MAR-2000 JP 2000087536

PF 27-MAR-2000 JP 2000087536

PF 27-MAR-2000 JP 200087536

PC C12N15/09,C07K14/415,C12N1/15,C12N1/19,C12N1/21,C12N5/10// PC

CC12N15/09,C12R1:91),(C12N5/10,C12R1:91),C12N15/00,C12R1:91)

CC Description of Artificial Sequence: Synthetic primer DNA FH

Location/Qualifiers.
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                                                                                                                                                                                                                                                                                      Peptide having a function
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        Location/Qualifiers.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Location/Qualifiers
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RESULT 488
BD089539/c
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BD056371
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Best Local Similarity
Matches 16; Conserv
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synthetic construct
artificial sequences.

E 1 (bases 1 to 20)

S Takagi,M., Shinji,H. and Ota,K.

S Tokagi,M., Shinji,H. and Ota,K.

L Patent: JP 2001269176-A 5 02-OCT-2001;

AGENCY OF IND SCIENCE & TECHNOL

OS Artificial Sequence
PN JP 2001269176-A/5

PD 02-OCT-2001
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                                                       A method of arraying genome clone Patent: JP 2001321190-A 1783 20-NOV-2001; THE INSTITUTE OF PHYSICAL AND CHEMICAL RE
                                                                                                                 synthetic construct artificial sequences.
1 (bases 1 to 20)
                                                                                                                                                                            BD089539.1 GI:22635149
JP 2001321190-A/1783.
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JP 2001269176-A/5.
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Similarity 84.2%;
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Description of Artificial Sequence: Synthetic primer DNA FH
Location/Qualifiers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-MAR-2000 JP 2000087536
MASARU TAKAGI,HIDEAKI SHINJI,KEN OTA
C12N15/09,C07K14/415,C12N1/15,C12N1/19,C12N1/21,C12N5/10// PC
Artificial Sequence
JP 2001321190-A/1783
20-NOV-2001
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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Pred. No. 2.5e+02;
0; Mismatches 3
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Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              regulating transcription of gene.
                                                         AND CHEMICAL RESEARCH, YUGENKAISHA
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RESULT 489
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AUTHORS
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Best Local Similarity
Matches 16; Conserv
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PI EIICHI
PC C12N15/00,
C12N15/00,
PC C12N15/
CC Descrip
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PD 09-JUL-2002
PF 27-DEC-2000 JP 2000399573
PF 27-DEC-2000 JP 2000399573
PI MICHIFUMI TANGA,HIROSHI OKAMURA,KOJIRO TAKAHASHI PC C12N15/09,C12N15/09,C12N15/00 CC Mei C12N15/09,C12N15/09,C12N15/00 CC Mei C12N15/09,C12N15/09 CC Mei C12N15/09 out thermal cycle of PCR using DNA- CC
                                                                                                                                                                                                                                                                                                                                                                                       Patent: JP 2002191369-A 1 09-JUL-2002;
TOYO KOHAN CO LTD, KOJIRO TAKAHASHI
OS Artificial Sequence
PN JP 2002191369-A/1
PD 09-JUL-2002
PF 27-DEC-2000 JP 2000399573
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1 (bases 1 to 20)

Tanga, M., Okamura, H. and Takahashi, K.
S47176S4 20 by lipoprotein lipase (introns 3, segment 4 of 5).
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BD161924.1 GI:27867682
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Method for carrying out thermal
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EIICHI SOEDA
C12N15/09,C12N15/09,C12M1/00,C12Q1/68,G01N33/53,G01N33/566,
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No. 2.5e+02;
0; Mismatches 3;
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Pred. No. 2.5e+02;
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8} [human, Genomic, 20 nt,
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1 (bases 1 to 20)

Gotoda, T., Yamada, N., Murase, T., Shimano, H., Shimada, M., Harada, K., Kawamura, M., Kozaki, K. and Yazaki, Y.

Detection of three separate DNA polymorphisms in the human lipoprotein lipase gene by gene amplification and restriction endonuclease digestion

J. Lipid Res. 33 (7), 1067-1072 (1992)
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                                                                                                                                                                        Submitted (04-AUG-2001) Akira Horii, Tohoku University School of Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, Sendai, Miyagi 980-8575, Japan (E-mall:horii@mall.cc.tohoku.ac.jp, Tel:81-22-717-8047)
                                                                                                                                                                                                                                                                                                                       A BAC-based STS-content map spanning a 35-Mb region of human chromosome 1p35-p36 Genomics 74 (1), 55-70 (2001) 21269192
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Watanabe,N., Inazawa,J., Hosoda,F., Arai,Y., Mizushima,H.,
Morohashi,A., Ohira,M., Nakagawara,A., Liu,S., Hoshi,M., Horii,A.
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/note="forward primer for human STS sts-H96854 at 1p36 sts-H96854 obtained from clones B293A18, B122E3, B91D18, Human BAC library RPCI-11"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                           /db_xref="taxon:32630"
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                                                                                              /organism="synthetic construct"
/mol_type="genomic DNA"
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0; Mismatches
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